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Tyr His Ile Ile His Arg Asp Ile Lys Pro Asp Asn Ile Leu Leu Asp
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Glu His Gly His Val His Ile Thr Asp Phe Asn Ile Ala Thr Val Val
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Cys Lys Gly Met Val Ala Leu Leu Arg Lys Leu Leu Thr Lys Asp Pro
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Glu Ser Arg Val Ser Ser Leu His Asp Ile Gln Ser Val Pro Tyr Leu
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Lys Met Asp Leu Pro Pro Gly Asp Pro Gly Val Leu Pro Leu Ser Cys
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Pro Gln Glu Cys Pro Asp Pro His Ser Tyr Pro Gly Pro Arg Ser Pro
                                                45
                            40
Thr Pro Gly Leu Pro Ser Ser Ala Val Asn Asp Asp Leu Leu Leu
Pro Ser Ser Leu Pro Ser Val Thr Lys Gly Leu Pro Arg Cys Gln Leu
                                         75
                    70
Trp Asn Glu Gly Cys Pro Trp Glu Val Met Ile Leu Arg Tyr Thr Gly
                                     90
Ala Gln Gln Ile Ala Ser Ser Tyr Pro Gln Thr Val Phe Ala Cys Met
                                 105
            100
Gln Pro Leu Ala Leu Pro Leu Cys Gly Arg Lys Pro Ala Gln Gly His
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Thr Ala Gly Gln Gln His Ser Trp Ser Gln Ile
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 gaactgggcc tagaacttgt ttttgtgtgg aaccgtgacc ctggacgaat ggcagggagt
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 gtgccccctg ccctgcagct cgaagacctc actacacttg aggaaaggca ccctgacctt
 300
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 gcaaaccttc tgagccttcg tgtcaccatg gccacacacc ccgatggctt ccggcttgag
 420
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ggacccctgg ctgcagccca cagccctggg ccttgcactg tgctctacga aggccctgtc
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Asp Arg Gly Pro Trp Arg Val Gly Val Val Gly Tyr Gly Arg Leu Gly
                            40
Gln Ser Leu Val Ser Arg Leu Leu Ala Gln Gly Ser Glu Leu Gly Leu
                        55
Glu Leu Val Phe Val Trp Asn Arg Asp Pro Gly Arg Met Ala Gly Ser
                                        75
Val Pro Pro Ala Leu Gln Leu Glu Asp Leu Thr Thr Leu Glu Glu Arg
                                    90
                85
His Pro Asp Leu Val Val Glu Val Ala His Pro Lys Ile Ile His Glu
                                105
Ser Gly Val Gln Ile Leu Arg His Ala Asn Leu Leu Ser Leu Arg Val
                            120
Thr Met Ala Thr His Pro Asp Gly Phe Arg Leu Glu Gly Pro Leu Ala
                        135
                                            140
Ala Ala His Ser Pro Gly Pro Cys Thr Val Leu Tyr Glu Gly Pro Val
                                        155
                    150
Arg Gly Leu Cys Pro Phe Ala Pro Arg Asn Ser Asn Thr Met Ala Ala
                                    170
Ala Ala Leu Ala Ala Pro Ser Leu Gly Phe Asp Gly Val Ile Gly Val
                                185
            180
Leu Val Ala Asp Thr Ser Leu Thr Asp Met His Val Val Asp Val Glu
                             200
Leu Ser Gly Pro Arg Gly Pro Thr Gly Arg Ser Phe Ala Val His Thr
                         215
Arg Arg Glu Asn Pro Ala Glu Pro Gly Ala Val Thr Gly Ser Ala Thr
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240
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225
Val Thr Ala Phe Trp Arg Ser Leu Leu Ala Cys Cys Gln Leu Pro Ser
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Arg Pro Gly Ile His Leu Cys
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tcaagcattg gaagaattta gggaaaaaaa tcagagatta caaaaattat gggttggaag
attaattotg tattootoag ttototatot gtttacatgo ttaattgtat atttgtggta
240
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 gacaaccagt ttaatgaaga atctttagaa cacgatgttc ttgatgataa tacagagcag
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 tcagaggaac cagaggagaa acaagagact gagaatgagg aagcctcagt gattgaaacc
 1260
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aactccacag ttcctggagc tgattctatt cctgatcctg aactaagtgg agaatctttg
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1365
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<211> 194
<212> PRT
<213> Homo sapiens
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Arg Glu Arg Gly Ala Leu Asp Arg Ile Val Glu Tyr Leu Val Gly Asp
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Gly Pro Gln Asn Arg Tyr Ala Leu Ile Cys Gln Gln Cys Phe Ser His
                                                45
                            40
Asn Gly Met Ala Leu Lys Glu Glu Phe Glu Tyr Ile Ala Phe Arg Cys
                        55
Ala Tyr Cys Phe Phe Leu Asn Pro Ala Arg Lys Thr Arg Pro Gln Ala
                                        75
                    70
Pro Arg Leu Pro Glu Phe Ser Phe Glu Lys Arg Gln Val Val Glu Gly
                                    90
                     •
                85
Ser Ser Ser Val Gly Pro Leu Pro Ser Gly Ser Val Leu Ser Ser Asp
                                105
Asn Gln Phe Asn Glu Glu Ser Leu Glu His Asp Val Leu Asp Asp Asn
                            120
 Thr Glu Gln Thr Asp Asp Lys Ile Pro Ala Thr Glu Gln Thr Asn Gln
                                            140
                        135
 Val Ile Glu Lys Ala Ser Asp Ser Glu Glu Pro Glu Glu Lys Gln Glu
                                         155
                    150
 Thr Glu Asn Glu Glu Ala Ser Val Ile Glu Thr Asn Ser Thr Val Pro
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 Gly Ala Asp Ser Ile Pro Asp Pro Glu Leu Ser Gly Glu Ser Leu Thr
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                                 185
 Ala Glu
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 <212> DNA
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 gaggeettee etgeecagte eccaeaggae etcaectagg gtggaggaga geaacageaa
 geteetggag teagagagga agetgeagga ggagegaeae egeaeegtgg tettggagea
  acatetggag aagataegee tggageeagg gaaggeatea geeteeeaga gageagetee
  300
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caggaccaaa acagctccgc tcctggatgt atgctgtgta cggggccttg gctgtgatgg
360
gcacaatggg cccttggtac ctgctgctgc tgcttggtca ctgtgtgggc ctctatgtgg
cetegetttt gggccagece tggctetgte ttggcettgg ettggecage etggeeteet
tcaagatgga ccccctaatc tcttggcaga gcgggtttgt aacaggcact tttgatcttc
aagaggtgct gtttcatggg ggcagcagct tcacagtgct gcgttgcacc agctttgcac
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gcctattcaa acctggtgta tgactgggtg aaggcggccg tcctctttgg tgttgtcaac
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tacgtctttg cggaaacgca ctttgaccgt ggcatcaacg actggctttg caaatatgtg
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 1320
 ggagccatga acttctgggc catcatcatg tacaaccttg tgagcctgaa cagcctcaaa
 1380
 ttcacagage tggttgeeeg gegeetgeta etcacagggt tececeagae caegetgtee
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 1500
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 1620
 ttctacaaca aaaaaaa
 1637
 <210> 4452
  <211> 328
  <212> PRT
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  <400> 4452
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Lys Tyr Asn Phe Tyr Leu Pro Phe Phe Phe Gly Pro Ile Met Thr
                           40
Phe Asp Arg Phe His Ala Gln Val Ser Gln Val Glu Pro Val Arg Arg
                                          60
                       55
Glu Gly Glu Leu Trp His Ile Arg Ala Gln Ala Gly Leu Ser Val Val
Ala Ile Met Ala Val Asp Ile Phe Phe His Phe Phe Tyr Ile Leu Thr
               85
Ile Pro Ser Asp Leu Lys Phe Ala Asn Arg Leu Pro Asp Ser Ala Leu
                              105
Ala Gly Leu Ala Tyr Ser Asn Leu Val Tyr Asp Trp Val Lys Ala Ala
                                              125
                          120
Val Leu Phe Gly Val Val Asn Thr Val Ala Cys Leu Asp His Leu Asp
                       135
Pro Pro Gln Pro Pro Lys Cys Ile Thr Ala Leu Tyr Val Phe Ala Glu
                   150
                                      155
Thr His Phe Asp Arg Gly Ile Asn Asp Trp Leu Cys Lys Tyr Val Tyr
                                  170
                165
Asn His Ile Gly Gly Glu His Ser Ala Val Ile Pro Glu Leu Ala Ala
                               185
            180
Thr Val Ala Thr Phe Ala Ile Thr Thr Leu Trp Leu Gly Pro Cys Asp
                                     205
                            200
Ile Val Tyr Leu Trp Ser Phe Leu Asn Cys Phe Gly Leu Asn Phe Glu
                        215
Leu Trp Met Gln Lys Leu Ala Glu Trp Gly Pro Leu Ala Arg Ile Glu
                                       235
                  230
Ala Ser Leu Ser Val Gln Met Ser Arg Arg Val Arg Ala Leu Phe Gly
                                   250
 Ala Met Asn Phe Trp Ala Ile Ile Met Tyr Asn Leu Val Ser Leu Asn
                                                   270
                                265
 Ser Leu Lys Phe Thr Glu Leu Val Ala Arg Arg Leu Leu Thr Gly
                           280
 Phe Pro Gln Thr Thr Leu Ser Ile Leu Phe Val Thr Tyr Cys Gly Val
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 Gln Leu Val Lys Glu Arg Glu Arg Thr Leu Ala Leu Glu Glu Glu Gln
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 Lys Gln Asp Lys Glu Lys Pro Glu
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 <212> DNA
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gcacatctat acceactctg gctctgaaag gcttgtcaac caaaaatggg cagctggggc

180

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taaggcatat ttaaacaaag gctccaaagg accectttca cttgggtcta gcatccagcc
240
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ccaaggcatg ccatcactgc agcactcaac cctctggtca cagtggagtc gccggtccag
cetgaaatat tactacagag gagaaagace cattettget atgttgetet atettecacg
tccaaaaaca gtcctatgta gcttcagctg ctccgaaatc aggtcacaga acagcaggag
acatteettt ggeaaaaaag gacaegettt tgteetgtat ettataetgg taagtgaage
tetgateceg gtggaetgeg ggetgegatg gteteeteea eaggateete agetacagag
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685
<210> 4454
<211> 207
<212> PRT
<213> Homo sapiens
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Met Ile Ile Leu Val Val Thr Leu His Thr Cys His Pro Val Pro Ser
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Pro Gly Trp His Ile Tyr Thr His Ser Gly Ser Glu Arg Leu Val Asn
Gln Lys Trp Ala Ala Gly Ala Lys Ala Tyr Leu Asn Lys Gly Ser Lys
                            40
Gly Pro Leu Ser Leu Gly Ser Ser Ile Gln Pro Leu Ser Gln Gln Arg
                        55
Gln Asp Cys Gly Pro Leu Cys Phe Leu Asn Arg Ala Gln Gly Ser Gln
                                         75
                    70
Gly Met Pro Ser Leu Gln His Ser Thr Leu Trp Ser Gln Trp Ser Arg
                                     90
Arg Ser Ser Leu Lys Tyr Tyr Tyr Arg Gly Glu Arg Pro Ile Leu Ala
                                                     110
                                 105
             100
 Met Leu Leu Tyr Leu Pro Arg Pro Lys Thr Val Leu Cys Ser Phe Ser
                             120
 Cys Ser Glu Ile Arg Ser Gln Asn Ser Arg Arg His Ser Phe Gly Lys
                         135
                                             140
 Lys Gly His Ala Phe Val Leu Tyr Leu Ile Leu Val Ser Glu Ala Leu
                                         155
                    150
 Ile Pro Val Asp Cys Gly Leu Arg Trp Ser Pro Pro Gln Asp Pro Gln
                                     170
                 165
 Leu Gln Arg Gln Arg Arg Met Lys Glu Glu Gln Pro Pro Gln Asp Leu
                                185
 Leu His Trp Glu Pro His Pro Thr Phe Ser Val Pro Phe Thr Arg
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<212> DNA

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105
            100
Lys Gln Gln Ser Glu Asp Asp Val Arg Arg Leu Phe Glu Ala Phe Gly
                            120
                                                125
Asn Ile Glu Glu Cys Thr Ile Leu Arg Gly Pro Asp Gly Asn Ser Lys
                                            140
                        135
Gly Cys Ala Phe Val Lys Tyr Ser Ser His Ala Glu Ala Gln Ala Ala
                    150
Ile Asn Ala Leu His Gly Ser Gln Thr Met Pro Gly Ala Ser Ser Ser
                                    170
                165
Leu Val Val Lys Phe Ala Asp Thr Asp Lys Glu Arg Thr Met Arg Arg
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                                185
            180
Met Gln Gln Met Ala Gly Gln Met Gly Met Phe Asn Pro Met Ala Ile
                            200
        195
Pro Phe Gly Ala Tyr Gly Ala Tyr Ala Gln Ala Leu Met Gln Gln Gln
                                            220
                        215
Ala Ala Leu Met Ala Ser Val Ala Gln Gly Gly Tyr Leu Asn Pro Met
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Ala Ala Phe Ala Ala Ala Gln Met Gln Met Ala Ala Leu Asn Met
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Asn Gly Leu Ala Ala
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 780
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1260
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1380
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 Gln Leu Leu Met Tyr Gln Gln His Thr Ser His Tyr Asp Leu Glu Arg
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             20
 Lys Gly Gly Tyr Leu Met Leu Ser Phe Ile Asp Phe Cys Pro Phe Ser
                             40
                                                 45
 Val Met Arg Leu Arg Ser Leu Pro Ser Pro Gln Arg Tyr Thr Arg Gln
 Glu Arg Tyr Arg Ala Arg Pro Pro Arg Val Leu Glu Arg Ser Gly Phe
                                         75
                     70
 His Asn Glu Asn Ser Leu Ala Ile Tyr Gln Gly Leu Val Tyr Tyr Leu
 Leu Trp Leu His Ser Val Tyr Asp Lys Asp Tyr Tyr Phe Phe Leu Ala
                                 105
             100
 Ser Asn Trp Arg Ser Ala Gly Gly Val Ser Ile Glu Met Asp Ser Tyr
                             120
         115
 Glu Lys Ile Tyr Asn Leu Glu Ser Ala Tyr Glu Leu Pro Glu Arg Ile
                                              140
                         135
 Phe Leu Asp Lys Gly Thr Glu Tyr Ser Phe Ala Ile Phe Leu Ser Ala
                                          155
                     150
 Gln Gly His Ser Phe Arg Thr Gln Ser Glu Leu Gly Leu Arg Gly Thr
                                      170
 Arg Val Glu Pro Glu Gly Arg Gly Glu Gly Tyr Gln Asn Leu Gly Ala
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190
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            180
Trp Gly Ala Gly Thr Pro Ser Glu Gly Arg Gly Leu Ser Val Asp Val
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Gly Val Val Leu Ala Asp Pro Gly Cys Ile Glu Ala Ser Val Lys Gln
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Glu Val Leu Ile Asn Arg Asn Ser Val Leu Phe Ser Ile Thr Leu Lys
                                        235
                   230
Asp Lys Lys Leu Cys Tyr Asp Gln Gly Ile Ser Gly His His Leu Met
                                    250
                245
Glu Thr Ser Met Thr Val Asn Val Arg Ser Lys Pro Gly Gly Glu Gly
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            260
Lys Arg Leu Ala Phe Asp Ile Thr Tyr Thr Leu Glu Tyr Ser Arg Leu
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                                                285
        275
Lys Asn Lys His Tyr Phe Asp Cys Val Asn Val Asn Pro Glu Met Pro
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Cys Phe Leu Phe Arg Asp Ser Val Tyr Val Leu Leu Val Val Gly Gly
                                        315
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Gly Pro Thr Leu Asp Ser Leu Lys Asp Tyr Ser Glu Asp Glu Ile Tyr
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Arg Phe Asn Ser Pro Leu Asp Lys Thr Asn Ser Leu Ile Trp Thr Thr
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Arg Thr Thr Arg Thr Thr Lys Asp Ser Ala Phe His Ile Met Ser His
                            360
Glu Ser Pro Gly Ile Glu Trp Leu Cys Leu Glu Asn Ala Pro Cys Tyr
                        375
                                            380
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                    390
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<211> 1114
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540
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 Ser Arg Gly Arg Ala Ala Asn Gly Arg Ala Pro Pro Gly Pro Leu Thr
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 Arg Arg Leu Ala Gly Arg Ala Arg Thr Pro Arg Pro Lys Trp Leu Phe
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Leu Ser Ile Lys Tyr Met Ala Arg Ser Phe Arg Gly Ala Val Ala Ile
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					gaaaccagct
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7 500					a gaagatcatt
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					t gtgacagggc
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 Arg Pro Lys Asp Pro Tyr Cys His Pro Val Cys Ala Asn Arg Phe Ser
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 Gly Val Leu Gly Thr Glu Ala His Ser Glu Val Thr Phe Asp Met Glu
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 Ile Leu Gly Ile Ile Ser Thr Ile Tyr Lys Phe Gln Gly Met Ser Asp
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 Phe Gln Tyr Leu Ala Val His Thr Glu Ala Gly Gly Lys His Thr Ser
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 Met Tyr Asp Lys Val Leu Met Leu Arg Pro Glu Lys Glu Ala Phe Phe
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 Asp Ala Pro Val Asp Tyr Phe Tyr Arg Pro Glu Thr Gln His Arg Glu
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 Gly Tyr Asn Asn Pro Pro Ile Ser Gly Glu Asn Leu Ile Gly Leu Ser
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200

195

205

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Leu Phe Asp Ile Arg Pro Ile Trp Ser Arg Asn Ala Val Lys Ala Asn
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Ile Ser Val His Pro Asp Lys Leu Lys Val Leu Leu Pro Phe Ile Ala
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Phe Arg Ile Arg Cys Gly Met Lys His Gly Tyr Ala Pro Ser Asp Leu
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Pro Val Lys Ala Lys Arg Ser Thr Tyr Asn Tyr Ser Leu Pro Ile Thr
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Val Lys Lys Thr Ser Ser Gln Leu Val Thr Met His Asp Leu Lys Gln
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Gly Leu Gly Arg Ser Gly Thr Ser Gly Ala Arg Lys Pro Ala Ser Ser
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Lys Tyr Lys Leu Lys Asp Ser Val Tyr Ile Phe Arg Glu Gly Ala Leu
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Pro Pro Tyr Arg Gln Met Phe Tyr Gln Leu Cys Asp Leu Asn Val Glu
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Glu Leu Gln Lys Ile Ile His Arg Asn Asp Gly Ala Glu Asn Ser Cys
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Asp Thr Met Ser Leu Met Ile Arg Gln Thr Ile Arg Ser Lys Arg Pro
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 Asp Thr Ile Gly Gln Met Arg Arg Xaa Ala Val Gly Leu Val Asp Ala
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 Val Lys Ala Thr Asp Gln Tyr Cys Ala Arg Leu Arg Gln Ala Gly Ser
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Phe Gly Glu Gly Leu Leu Glu Ala Glu Leu Ala Ala Leu Cys Pro Thr
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Thr Leu Ala Pro Tyr Tyr Leu Arg Ala Pro Ser Val Ala Leu Pro Val
                                             60
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Ala Gln Val Pro Thr Asp Pro Gly His Phe Ser Val Leu Leu Asp Val
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Lys His Phe Ser Pro Glu Glu Ile Ala Val Lys Val Val Gly Glu His
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 Val Glu Val His Ala Arg His Glu Glu Arg Pro Asp Glu His Gly Phe
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 Val Ala Arg Glu Phe His Arg Arg Tyr Arg Leu Pro Pro Gly Val Asp
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Cys Pro Asp Asp Thr His Phe Val Ser Ser Ser Ser Asp Lys Ser Val
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Ser					٥.5					90					95	
Asn Thr Cys Lys Asp Asn Pro Cys Gly Arg Gly Gln Cys Leu Ile Thr  115			<b>.</b> _	<b>+</b>	85	D~C	Dhe	Ser			Lvs	Cvs	Gln			Gln
Asn Thr Cys Lys Asp Asn Pro Cys Gly Arg Gly Gln Cys Leu Ile Thr	Cys S	er			Ala	PLO	FIIC	261	105			-1-		110		
115	m	N /	~	100	Acn	Δen	Pro	Cvs	Glv	Arq	Gly	Gln	Cys	Leu	Ile	Thr
Gln Ser Pro Pro Tyr Tyr Arg Cys Val Cys Lys His Pro Tyr Trr Gly 130 130 135 150 150 155 160 155 160 175 165 160 175 165 160 175 165 160 175 165 160 175 165 160 175 165 170 175 180 180 180 180 180 185 190 180 185 190 180 180 195 190 200 205 205 207 207 215 210 210 215 210 225 230 235 240 240 241 241 241 241 241 242 243 244 245 240 241 242 241 242 242 244 245 240 241 242 244 245 240 241 242 244 245 244 245 244 245 244 245 245			115					120					125			
130	G1 G	ا مد	D.C.O.	Dro	Tyr	Tvr	Ara	Cvs	Val	Cys	Lys	His	Pro	Tyr	Thr	Gly
Pro Ser Cys Ser   Gln   Val   Val   Pro   Val   Cys   Arg   Pro   Asn   Pro   Cys   Gln   145		20					135					140				
150	D~~ C	.30	Cve	Ser	Gln	Val	Val	Pro	Val	Сув	Arg	Pro	Asn	Pro	Cys	Gln
Asn Gly Ala Thr Cys Ser Arg His Lys Arg Arg Ser Lys Phe Try Cys 165	7.45					150					155					100
165	Yen C	ilv.	Δla	Thr	Cvs	Ser	Arg	His	Lys	Arg	Arg	Ser	Lys	Phe	Thr	Cys
Ala Cys Pro Asp Gln Phe Lys Gly Lys Phe Cys Glu I1e Gly Ser Asp 180					165					170					1/3	
Asp Cys Tyr Val Gly Asp Gly Tyr Ser Tyr Arg Gly Lys Met Asn Arg  195 200 205  Thr Val Asn Gln His Ala Cys Leu Tyr Trp Asn Ser His Leu Leu Leu 210 215 220 230 230 230 230 230 230 230 230 230	Δla (	'vs	Pro	Asp	Gln	Phe	Lys	Gly	Lys	Phe	Cys	Glu	Ile	Gly	Ser	Asp
Thr Val Asn Gln His Ala Cys Leu Tyr Trp Asn Ser His Leu Leu Leu 210	•			100					185					130		
Thr Val Asn Gln His Ala Cys Leu Tyr Trp Asn Ser His Leu Leu Leu 210	Asp (	Cvs	Tyr	Val	Gly	Asp	Gly	Tyr	Ser	Tyr	Arg	Gly	Lys	Met	Asn	Arg
Simple   S			105					200					∠∪5			
Simple   S	Thr V	Val	Asn	Gln	His	Ala	Cys	Leu	Tyr	Trp	Asn	Ser	His	Leu	Leu	Leu
225		210					215					220				
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Cys Phe Ile Lys Val Thr Asn Asp Lys Val Lys Trp Glu Tyr Cys Asp 260	225					230					235					240
Cys Phe Ile Lys Val Thr Asn Asp Lys Val Lys Trp Glu Tyr Cys Asp 260	Gly (	Glu	His	Asn	Phe	Cys	Arg	Asn	Pro	Asp	Ala	Asp	GIu	гуѕ	PIO	пр
Val Ser Ala Cys Ser Ala Gln Asp Val Ala Tyr Pro Glu Glu Ser Pro 275  280  Thr Glu Pro Ser Thr Lys Leu Pro Gly Phe Asp Ser Cys Gly Lys Thr 290  Glu Ile Ala Glu Arg Lys Ile Lys Arg Ile Tyr Gly Gly Phe Lys Ser 300  Thr Ala Gly Lys His Pro Trp Gln Ala Ser Leu Gln Ser Ser Leu Pro 325  Leu Thr Ile Ser Met Pro Gln Gly His Phe Cys Gly Gly Ala Leu Ile 340  His Pro Cys Trp Val Leu Thr Ala Ala His Cys Thr Asp Ile Lys Thr 355  Arg His Leu Lys Val Val Leu Gly Asp Gln Asp Leu Lys Lys Glu Glu 370  Phe His Glu Gln Ser Phe Arg Val Glu Lys Ile Phe Lys Tyr Ser His 385  Tyr Asn Glu Arg Asp Glu Ile Pro His Asn Asp Ile Ala Leu Leu Lys 400  Thr Val Cys Leu Pro Asp Gly Ser Phe Pro Ser Gly Ser Glu Cys His 435  Ile Ser Gly Trp Gly Val Thr Glu Thr Glu Thr Gly Lys Gly Ser Arg Gln Leu 450  Leu Asp Ala Lys Val Lys Leu Ile Ala Asn Thr Leu Cys Asn Ser Arg 465  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 500  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 510					245					250					2,7,5	
Val Ser Ala Cys Ser Ala Gln Asp Val Ala Tyr Pro Glu Glu Ser Pro 275	Cys	Phe	Ile	Lys	Val	Thr	Asn	Asp	Lys	Val	Lys	Trp	GIU	171	Cys	ASP
Thr Glu Pro Ser Thr Lys Leu Pro Gly Phe Asp Ser Cys Gly Lys Thr 290  Glu Ile Ala Glu Arg Lys Ile Lys Arg Ile Tyr Gly Gly Phe Lys Ser 330  Thr Ala Gly Lys His Pro Trp Gln Ala Ser Leu Gln Ser Ser Leu Pro 335  Leu Thr Ile Ser Met Pro Gln Gly His Phe Cys Gly Gly Ala Leu Ile 340  His Pro Cys Trp Val Leu Thr Ala Ala His Cys Thr Asp Ile Lys Thr 355  Arg His Leu Lys Val Val Leu Gly Asp Gln Asp Leu Lys Lys Glu Glu 370  Phe His Glu Gln Ser Phe Arg Val Glu Lys Ile Phe Lys Tyr Ser His 385  Tyr Asn Glu Arg Asp Glu Ile Pro His Asn Asp Ile Ala Leu Leu Lys 405  Leu Lys Pro Val Asp Gly His Cys Ala Leu Glu Ser Lys Tyr Val Lys 420  Thr Val Cys Leu Pro Asp Gly Ser Phe Pro Ser Gly Ser Glu Cys His 450  Leu Asp Ala Lys Val Lys Leu Ile Ala Asn Thr Leu Cys Asn Ser Arg 465  Leu Asp Ala Lys Val Lys Leu Ile Asp Asp Ser Met Ile Cys Asn Ser Arg 485  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro Ser Gly Ser Gly Ser Arg Gln Leu Asp Ala Lys Val Dys A85  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro Ser Gly Ser Arg Gln Leu Asp Ala Lys Val Lys Leu Ile Ala Asn Thr Leu Cys Asn Ser Arg A85  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro Ser Gly Gly Pro Ser Gly Gly Pro Ser Gln Leu Asp Ala Lys Val Lys Leu Ile Ala Asn Thr Leu Cys Asn Ser Arg A85  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro Ser Gly Gly Pro Ser Gln Lys Pro Gly Gln Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro Ser Gly Gly Pro Ser Gln Cys File Cys Asn Ser A85  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro Ser Gly Gly Pro Ser Gln Cys File Cys Asn Ser A85  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro Ser Glo Cys File Cys Gln Gly Asp Ser Gly Gly Pro Ser Gly Gly Pro Ser Glo Cys File Cys Gly Gly Pro Ser G				260					265					210		
Thr Glu Pro Ser Thr Lys Leu Pro Gly Phe Asp Ser Cys Gly Lys Thr 290	Val	Ser	Ala	Cys	Ser	Ala	Gln	Asp	Val	Ala	Tyr	Pro	205	Gru	261	110
290			275				_	280	<b>~1</b>	Db =	7.00	Cor		Glv	Lvs	Thr
Glu Ile Ala Glu Arg Lys Ile Lys Arg Ile Tyr Gly Gly Phe Lys Ser 310	Thr	Glu	Pro	Ser	Thr	Lys	Leu	Pro	GIĀ	Pne	Asp	300	Суз	GIY	шуБ	
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Thr Ala Gly Lys His Fro Trp Gln Ala Ser Leu Gln Ser Ser Leu Pro 325  Leu Thr Ile Ser Met Pro Gln Gly His Phe Cys Gly Gly Ala Leu Ile 340  His Pro Cys Trp Val Leu Thr Ala Ala His Cys Thr Asp Ile Lys Thr 350  Arg His Leu Lys Val Val Leu Gly Asp Gln Asp Leu Lys Lys Glu Glu 370  Phe His Glu Gln Ser Phe Arg Val Glu Lys Ile Phe Lys Tyr Ser His 385  Tyr Asn Glu Arg Asp Glu Ile Pro His Asn Asp Ile Ala Leu Leu Lys 405  Leu Lys Pro Val Asp Gly His Cys Ala Leu Glu Ser Lys Tyr Val Lys 420  Thr Val Cys Leu Pro Asp Gly Ser Phe Pro Ser Gly Ser Glu Cys His 435  Ile Ser Gly Trp Gly Val Thr Glu Thr Gly Lys Gly Ser Arg Gln Leu Asp Ala Lys Val Lys Leu Ile Ala Asn Thr Leu Cys Asn Ser Arg 465  Gln Leu Tyr Asp His Met Ile Asp Asp Ser Met Ile Cys Ala Gly Asn 485  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 505  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 505	Glu	Ile	Ala	Glu	Arg	Lys	IIe	ьуs	Arg	TIE	215	GLY	GLY			320
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Leu Thr Ile Ser Met Pro Gln Gly His Phe Cys Gly Gly Ala Leu Ile 340	Thr	Ala	Gly	Lys			Trp	6111	Ата	351	1				335	
His Pro Cys Trp Val Leu Thr Ala Ala His Cys Thr Asp Ile Lys Thr 355  Arg His Leu Lys Val Val Leu Gly Asp Gln Asp Leu Lys Lys Glu Glu 370  Phe His Glu Gln Ser Phe Arg Val Glu Lys Ile Phe Lys Tyr Ser His 385  Tyr Asn Glu Arg Asp Glu Ile Pro His Asn Asp Ile Ala Leu Lys Lys 400  Thr Val Cys Leu Pro Asp Gly His Cys Ala Leu Glu Ser Lys Tyr Val Lys 430  Thr Val Cys Leu Pro Asp Gly Ser Phe Pro Ser Gly Ser Glu Cys His 435  Leu Asp Ala Lys Val Lys Leu Ile Ala Asn Thr Leu Cys Asn Ser Arg Gln Leu Asp Ala Leu Tyr Asp His Met Ile Asp Asp Ser Met Ile Cys Ala Gly Asn 485  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 500  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 500				_	325	D===	~1 n	. Cla	uic	Dhe	· · Cvs	Glv	Glv	Ala		
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Arg His Leu Lys Val Val Leu Gly Asp Gln Asp Leu Lys Lys Glu Glu 370		_		340	) 	Τ Ο 11	Thr	· 101 =	Δla	His	. Cvs	Thr	Asp	Ile	Lys	Thr
Arg His Leu Lys Val Val Leu Gly Asp Gln Asp Leu Lys Lys Glu Glu 370	His	Pro			) vai	Leu	. 1111	360	1		,-		365	5	_	
Note		***	355	) . T.re	. 17-1	17 a 1	Ler	Gly	, Asr	Glr	ı Ast	Leu	Lys	Lys	: Glu	Glu
Phe         His         Glu         Gln         Ser         Phe         Arg         Val         Glu         Lys         Ile         Phe         Lys         Tyr         Ser         His           385         390         400         395         400         400         400         400         400         400         400         400         400         400         400         400         410         410         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         415         420         420         420         430         445         445         445         445         445         445         445         445         446         446         446		227					375	•				360	,			
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Leu Lys Pro Val Asp Gly His Cys Ala Leu Glu Ser Lys Tyr Val Lys 420	202	Nen	Gli	ι Δτο	r Ast	Glu	ı Ile	e Pro	His	. Ası	n Ası	o Ile	e Ala	ı Lev	ı Lev	ı Lys
Leu Lys       Pro Val Asp Gly His Cys       Ala Leu Glu Ser Lys       Tyr Val Lys         420       425       430         Thr Val Cys       Leu Pro Asp Gly Ser Phe Pro Ser Gly Ser Glu Cys His         435       440         1le Ser Gly Trp Gly Val Thr Glu Thr Gly Lys Gly Ser Arg Gln Leu         450       455         Leu Asp Ala Lys Val Lys Leu Ile Ala Asn Thr Leu Cys Asn Ser Arg         465       470         Gln Leu Tyr Asp His Met Ile Asp Asp Ser Met Ile Cys Ala Gly Asn         485       490         Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro         500					405					41(	U					,
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11e   Ser   Gly   Trp   Gly   Val   Thr   Glu   Thr   Gly   Lys   Gly   Ser   Arg   Gln   Leu     450	Thr	Val	Cv	s Le	u Pro	Ası	Gly	y Sei	c Phe	e Pro	o Se	r Gly	y Se	r Gli	ı Cys	s His
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465 470 475  Gln Leu Tyr Asp His Met Ile Asp Asp Ser Met Ile Cys Ala Gly Asn 485 490 495  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 500 500 510	Leu	Ast	Al	a Ly	s Val	l Lys	s Le	u Il	e Ala	a As	n Th	r Le	u Cy	s Ası	n Se	r Arg
Gln Leu Tyr Asp His Met Ile Asp Asp Ser Met Ile Cys Ala Gly Ash 485 490 495 Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 500 500 510	100					471	0				47	5				-100
485  Leu Gln Lys Pro Gly Gln Asp Thr Cys Gln Gly Asp Ser Gly Gly Pro 500  500  500  500	Gln	Let	ту	r As	p His	s Met	t Il	e As	p As	p Se	r Me	t Il	е Су	s Ala	a Gl	y Asn
505					4 9	Ε.				49	U				42	_
505	Leu	Glr	ı Ly	s Pr	o Gl	y Gl:	n As	p Th	r Cy	s Gl	n Gl	y As	p Se	r Gl	λ GT.	y Pro
Leu Thr Cys Glu Lys Asp Gly Thr Tyr Tyr Val Tyr Gly 11e Val Ser				50	Λ.				50	5				21	· ·	
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	Car	Δla	Leu	Glu	Lvs	Thr	Thr			Lys	Thr	Val	Glu	Ala	Pro	Leu
	Ser	125		. 014			125		-	-		126	0			
	Va1	Thr	Glu	Glu	Lvs	Thr	Val	Glu	Pro	Ala	Thr	Val	Ser	Glu	Glu	Ala
	126	5				127	0				127	5				1280
	Lvs	Pro	Ala	Ser	Glu	Pro	Ala	Pro	Ala	Pro	Val	Glu	Glr	Leu	Glu	Gln
					128	5				129	0				129	5
	Val	Asp	Leu	Pro	Pro	Gly	Ala	Asp	Pro	Asp	) Lys	Glu	. Ala	ı Ala	Met	Met
				130	0				130	5				131	.0	
	Pro	Ala	Gly	v Val	. Glu	ı Glu	Gly	Ser	Ser	Gly	/ Asp	Gln	Pro	Pro	Tyr	Leu
			131	.5				132	0				132	25		
	Asp	Ala	Lys	Pro	Pro	Thr			Ala	Ser	: Phe	Ser	Glr	ı Ala	Glu	Ser
	_	133	0				133	5				134	10			
	Asn	val	Ast	Pro	Gli	ı Pro	Asp	Ser	Thr	Glr	n Pro	Leu	ı Sei	: Lys	Pro	Ala
	LADI			-								_				
	134	5				135	50				135	55				1360
	134	5				135	50				135	55				1360 Ala

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	1365	;		1370			1375	
Thr Ala Asp	Ala Glu	Pro Asp	Ala As	n Gln	Lys Ala	Glu Al	a Ala	Pro
	1380		13	885		13	90	
Glu Ser Gln	Pro Pro	Ala Ser	Glu As	p Leu	Glu Val	Asp Pr	o Pro	Val
1395	i		1400			1405		
Ala Ala Lys	Asp Lys	Lys Pro	Asn Ly	s Ser	Lys Arg	Ser Ly	s Thr	Pro
1410		1419			1420			
Val Gln Ala	Ala Ala	Val Ser	Ile Va			Val Th	r Arg	
1425		1430			1435		_	1440
Ser Glu Arg	Ile Asp	Arg Glu	Lys Le			Asn Se		
	1445		_	1450		- 2 - 22	1455	
Gly Glu Ala		Leu Leu			Met Glu	Ala Gi	u Lys	шe
	1460			165			70	<b>C1.</b>
Thr Arg Thr		Lys Asn		la Ala	Asp Leu	1485	S PLO	Giu
1475	) D	0 3 war	1480	ar Ara	Ara Asn		a Ser	Val
Pro Ser Leu	Pro Leu			rg Arg	1500		9 501	Vai
1490 Tyr Ala Thr	Mah (1)	149!		an Ara			s Glu	Pro
=	Met Giy	1510	GIU A		1515	var by	5 014	1520
1505 Val Glu Gln	Dro Ara		Ara Is			Ara Gl	u Leu	
vai Giu Gin	1525		Arg D	1530		5 0-	1535	
Glu Ala Ala			Thr Di			Arg Pr		
GIU AIA AIA	1540	110 1111		545			50	•
Thr Arg Arg	Arg Ala	Asp Glu			Asn Glu	Ala Ly	s Glu	Pro
1555		nop cru	1560			1565		
Ala Glu Thr	Leu Lvs	Pro Pro		ly Trp	Arg Ser	Pro Ar	g Ser	Gln
1570		157			158			
Lys Thr Ala	Ala Gly	Gly Gly	Pro G	ln Gly	Lys Lys	Gly Ly	rs Asn	Glu
1585		1590			1595			1600
Pro Lys Val	Asp Ala	Thr Arg	Pro G	lu Ala	Thr Thr	Glu Va	l Gly	Pro
	1609			1610			1619	
Gln Ile Gly	Val Lys	Glu Ser			Pro Lys	Ala Al	.a Glu	Glu
	1620			625			30	
Glu Ala Gly	Ser Glu	Gln Lys		sp Arg	Lys Asp	Ala Gi	y Thr	Asp
1635	5	_, _,	1640	1 ~1	**-1 **-1	1645		Dro
Lys Asn Pro	Pro Glu			al Glu			тѕ цуѕ	PIO
1650	_	165		7	166		a yen	Ser
Ala Pro Glu	Lys Asn		Ser L	ys Arg	1675	SEL AL	.g Asii	1680
1665 Arg Leu Ala	wal han	1670	71⇒ S	er Leu		Val As	n Ala	
Arg Leu Ala	168		Ala 5	1690		Val 11	169!	5
Val Ser Pro			Ala G			Arg G		
val ser Pro	1700	ALC ALC		705	017 014		710	- 1
Val Val Ala		Pro Glu			Ser Pro	Gln Ly	s Glu	Asp
171		110 014	1720	02 024		1725		-
Gly Leu Ser	Ser Gln	Leu Lvs		sp Pro	Val Asp	Pro As	sp Lys	Glu
1730	501 0211	173		- <u>-</u>	174			
Pro Glu Lys	Glu Asp			er Gly	Pro Ser	Pro G	lu Ala	Thr
1745		1750		-	1755			1760
Gln Leu Ala	Lys Gln		Leu G	lu Gln	Ala Val	Glu H	is Ile	Ala
	176		_	1770			177	
Lys Leu Ala			Ser A	la Ala	Tyr Lys	Ala A	sp Ala	Pro
_	1780		1	785		1	790	
Glu Gly Leu		Glu Asp	Arg A	sp Lys	Pro Ala	His G	in Ala	Ser
4		_		_				

		1795	;				1800	þ				1805	;		
Glu	Thr			Ala	Ala	Ala			Ser	Ile	Ile			Ile	Ser
024	1810					1815		•			1820		_		
Glv			Glu	Asn	Phe	Pro	Ala	Pro	Pro	Pro	Tyr	Pro	Gly	Glu	Ser
1825					1830					1835					1840
Gln	Thr	Asp	Leu	Gln	Pro	Pro	Ala	Gly	Ala	Gln	Ala	Leu	Gln	Pro	Ser
•		-		1845					1850					1855	
Glu	Glu	Glv	Met	Glu	Thr	Asp	Glu	Ala	Val	Ser	Gly	Ile	Leu	Glu	Thr
		-	1860			-		1865					1870		
Glu	Ala	Ala	Thr	Glu	Ser	Ser	Arg	Pro	Pro	Val	Asn	Ala	Pro	Asp	Pro
_		1875					1880					1885			
Ser	Ala	Gly	Pro	Thr	Asp	Thr	Lys	Glu	Ala	Arg	Gly	Asn	Ser	Ser	Glu
	1890				_	1895					1900				
Thr	Ser	His	Ser	Val	Pro	Glu	Ala	Lys	Gly	Ser	Lys	Glu	Val	Glu	Val
1905					1910					1915					1920
Thr	Leu	Val	Arg	Lys	Asp	Lys	Gly	Arg	Gln	Lys	Thr	Thr	Arg	Ser	Arg
			-	1925					1930					1935	
Arq	Lys	Arg	Asn	Thr	Asn	Lys	Lys	Val	Val	Ala	Pro	Val	Glu	Ser	His
	-	_	1940					1945					1950		
Val	Pro	Glu	Ser	Asn	Gln	Ala	Gln	Gly	Glu	Ser	Pro	Ala	Ala	Asn	Glu
		1955					1960					1965			
Gly	Thr	Thr	Val	Gln	His	Pro	Glu	Ala	Pro	Gln	Glu	Glu	Lys	Gln	Ser
-	1970					1975					1980				
Glu	Lys	Pro	His	Ser	Thr	Pro	Pro	Gln	Ser	Cys	Thr	Ser	Asp	Leu	Ser
1989					1990					1995					2000
Lys	Ile	Pro	Ser	Thr	Glu	Asn	Ser	Ser	Gln	Glu	Ile	Ser	Val	Glu	Glu
				2005					2010					201	
Arg	Thr	Pro	Thr	Lys	Ala	ser	Val	Pro	Pro	Asp	Leu	Pro	Pro	Pro	Pro
			2020					2025					2030		
Gln	Pro	Ala	Pro	Val	Asp	Glu	Glu	Pro	Gln	Ala	Arg			Val	His
		2035					2040					2045			
Ser	Ile	Ile	Glu	Ser	Asp	Pro		Thr	Pro	Pro			Pro	Ser	Ile
	2050					2055					206			_	_
Pro	Ile	Pro	Thr	Leu		Ser	Val	Thr	Ala			Leu	Ser	Pro	
206					207					207			_		2080
Val	Ala	Ser	Gly			Pro	His	Gln			Pro	Thr	Lys		
		_	_	208				_	209		~ 7	_	<b></b> 1	209	
Glu	Trp	Ile		_	Gln	Glu	Glu			Ala	Gin	Ser			ser
		_	210		_			210		<b>3</b>	77 7	*	2110		Com
Pro	Ala			Pro	Asp	Thr			ser	Asp	vai			ser	Ser
		2115					2120		<b>D</b>	T	m	212		ח ד ת	The
Ser			Arg	Lys	Ile	Leu		Asp	Pro	ьуѕ			ser	ALA	Inr
_	2130			<b></b>	<b>6</b>	213		m1	<b>3</b> 7 -	<b>-</b> 1.	214		Dvc	val	Car
		Thr	Ser	Tnr		Val	Thr	Thr	Ата			GIU	PIO	vaı	2160
214		D	<b>G</b>	T	215		77.	Dwa	Dwo	215		17-1	λen	Car	
		Pro	Cys	Leu		Glu	Ala	PIO	Pro			vai	Asp	Ser	пуз
216		<b>T</b>	~1	<b>~1</b>	217		27~	Dwo	Dxo	217		λen	Λen	Sar	Glu
ьys	rro	ьeu			гÀS	Thr	ATG			val	TIIT	ASIL	219		GIU
<b>-</b>	<b>01</b>	<b>7.7</b> -	218		17-7	T	37 T	218		n ~~	Tara	G1			בומ
тте	GIN			GIU	val	Leu			HT 9	Asp	пур	220		val	A.a
D		219	_				220								
	T7 7			D	T	T1 -	TT 1	C	37-7	T7~	Car	D ~~	Mot	Dva	\/ <b></b>
PIO		Ile		Pro	Lys	Ile		Ser	Val	Ile			Met	Pro	Val
	221	Ile O	Ala			Ile 221 Ser	5				222	0			

2225	2230	22	35	2240
Pro Gln Thr Leu T		Ser Ala Le	u Thr Gly Leu	Val Asn
	245	2250	_	2255
Val Ser Leu Val P	ro Val Asn Al	a Leu Lys Gl	y Pro Val Lys	Gly Ser
2260		2265	227	0
Val Thr Thr Leu L	ys Ser Leu Va	L Ser Thr Pr	o Ala Gly Pro	Val Asn
2275	22	30	2285	
Val Leu Lys Gly P	ro Val Asn Va	l Leu Thr Gl	y Pro Val Asn	Val Leu
2290	2295		2300	
Thr Thr Pro Val A	sn Ala Thr Va	L Gly Thr Va	l Asn Ala Ala	Pro Gly
2305	2310		:15	2320
Thr Val Asn Ala A	ala Ala Ser Al	a Val Asn Al	a Thr Ala Ser	
	2325	2330		2335
Thr Val Thr Ala G	ly Ala Val Th			
2340		2345	235	
Thr Thr Gly Thr V				Ser Thr
2355	23		2365	77.' - D
Lys Cys Lys Gln A		a Asn Glu As		HIS Pro
2370	2375		2380	Com Clar
Gly Ser Met Pro V				2400
2385	2390		395 Nalionian	
Ala Gly Leu Arg V		r Giu Giy və 2410	i vai Leu Leu	2415
Ser Gly Gln Lys T	2405		a Sar Mla Ive	
ser GIY GIN LYS 1	.III GIU GIY FI	2425	243	
Gln Ile Pro Pro A	Ala Ser Ala Me			
2435	24		2445	
Ser Lys Ser Gln V				Pro Pro
2450	2455		2460	
Ser Lys Gly Pro G		a Gly Tyr Al	la Asn Val Ala	Thr His
2465	2470		175	2480
Ser Thr Leu Val I	Leu Thr Ala Gl	n Thr Tyr As	sn Ala Ser Pro	Val Ile
2	2485	2490		2495
Ser Ser Val Lys A	Ala Asp Arg Pr	o Ser Leu Gl	lu Lys Pro Glu	Pro Ile
2500		2505	251	0
His Leu Ser Val S	Ser Thr Pro Va	l Thr Gln Gl		Lys Val
2515		20	2525	
Leu Thr Gln Gly I		o Pro Val Le		Gln Leu
2530	2535		2540	
Val Leu Thr Pro S				Asp Pro
2545	2550		555	2560
Val Thr Leu Lys I			in Pro Ala Asn	
	2565	2570	D	2575
Ser Thr Leu Thr E	Pro His His Pr			
2580		2585	259	
Thr Glu Val Asn F				Asp Arg
2595		00 - T T	2605	Dro Ara
Thr Val Ser His I		а гув Leu As	sp Ala His Ser 2620	FIO ALY
2610	2615	w Dha Dwa A		Pro Ser
Pro Ser Gly Pro C			rg Ala Ser His 535	2640
2625 Ser Thr Ala Ser 3	2630 The Ala Leu Se			
		2650	ra im var ricc	2655
Ala Gly Ile Pro V	2645 Val Pro Glo Ph		er Tle His Pro	
	voir ero Gill Pl	C TTE DET D		

266	0			2665	5				2670	)	
Ser Val Ile Met						Gln	Thr	Val	Ser	T.e.11	Ser
2675	PIO PIO		2680		1111	GIII	1111	2685		Deu	DCI
His Leu Ser Gln	Gly Glu				Δen	Thr	Pro	Thr	T.e.13	Pro	Ser
	Gry Gru	2699		1100	71011		2700				001
2690					_				_		_
Ile Thr Tyr Ser			GIU	Ala	Leu			Pro	Arg	Ala	
2705	271					271					2720
Leu Gln Pro Gln	Gln Ile	Glu	Val	Arg	Ala	Pro	Gln	Arg	Ala	Ser	Thr
	2725				2730	)				2735	5
Pro Gln Pro Ala	Pro Ala	Glv	Val	Pro	Ala	Leu	Ala	Ser	Gln	His	Pro
274				2745					2750		
Pro Glu Glu Glu		Тэ гээ	uic			Val	בות	λνα			Δl =
	vai mis	_			110	var	AIU	2765			niu
2755			2760		~ 7	_					•••
Pro Val Gln Ser	Glu Val			Met	Gin	Ser			Arg	Leu	HIS
2770		2775					2780				
Pro Tyr Thr Val	Pro Arg	Asp	Val	Arg	Ile	Met	Val	His	Pro	His	Val
2785	279	0				2799	5				2800
Thr Ala Val Ser	Glu Gln	Pro	Arq	Ala	Ala	Asp	Gly	Val	Val	Lys	Val
	2805		_		2810	_	-			2819	
Pro Pro Ala Ser		Pro	Gln				T.VS	Glu	Δla		
	=	FIO	GIII	2825		Gry	шүз	014	2830		Lys
282			_								
Thr Pro Asp Ala	Lys Ala				Pro	Thr	Pro			vai	Pro
2835			2840					2845			
Val Pro Val Pro	Leu Pro	Ala	Pro	Ala	Pro	Ala	Pro	His	Gly	Glu	Ala
2850		2859	5				2860	)			
Arg Ile Leu Thr	Val Thr	Pro	Ser	Asn	Gln	Leu	Gln	Gly	Leu	Pro	Leu
=	287					2879		•			2880
2865	2071	J				40/3					
2865 The Pro Pro Val			His	Glv	Val			Va 1	His	Ser	
Thr Pro Pro Val	Val Val	Thr				Gln		Val	His		Ser
Thr Pro Pro Val	Val Val 2885	Thr			2890	Gln )	Ile			2895	Ser 5
Thr Pro Pro Val	Val Val 2885 Gln Glu	Thr		Tyr	2890 Gly	Gln ) Asp	Ile Ile		Thr	2899 Tyr	Ser 5
Thr Pro Pro Val Gly Glu Leu Phe 290	Val Val 2885 Gln Glu 0	Thr Tyr	Arg	Tyr 2905	2890 Gly	Gln ) Asp	Ile Ile	Arg	Thr 2910	2 <b>8</b> 95 Tyr	Ser His
Thr Pro Pro Val	Val Val 2885 Gln Glu 0	Thr Tyr	Arg Thr	Tyr 2905 Gln	2890 Gly	Gln ) Asp	Ile Ile	Arg Ala	Thr 2910 Ser	2 <b>8</b> 95 Tyr	Ser His
Thr Pro Pro Val Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915	Val Val 2885 Gln Glu O Leu Thr	Thr Tyr His	Arg Thr 2920	Tyr 2905 Gln	2890 Gly Dhe	Gln ) Asp Pro	Ile Ile Ala	Arg Ala 2925	Thr 2910 Ser	2899 Tyr ) Ser	Ser His
Thr Pro Pro Val Gly Glu Leu Phe 290 Pro Pro Ala Gln	Val Val 2885 Gln Glu O Leu Thr	Thr Tyr His	Arg Thr 2920	Tyr 2905 Gln	2890 Gly Dhe	Gln ) Asp Pro	Ile Ile Ala	Arg Ala 2925	Thr 2910 Ser	2899 Tyr ) Ser	Ser His
Thr Pro Pro Val Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915	Val Val 2885 Gln Glu O Leu Thr	Thr Tyr His Lys	Arg Thr 2920	Tyr 2905 Gln	2890 Gly Dhe	Gln ) Asp Pro	Ile Ile Ala	Arg Ala 2925 Pro	Thr 2910 Ser	2899 Tyr ) Ser	Ser His
Thr Pro Pro Val  Gly Glu Leu Phe 290  Pro Pro Ala Gln 2915  Gly Leu Pro Ser 2930	Val Val 2885 Gln Glu O Leu Thr	Thr Tyr His Lys 2935	Arg Thr 2920 Thr	Tyr 2905 Gln O Ala	2890 Gly Phe Ala	Gln ) Asp Pro Gln	Ile Ile Ala Gly 2940	Arg Ala 2925 Pro	Thr 2910 Ser Pro	2899 Tyr ) Ser Pro	Ser His Val
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu	Val Val 2885 Gln Glu O Leu Thr Arg Thr	Thr Tyr His Lys 2935 Pro	Arg Thr 2920 Thr	Tyr 2905 Gln O Ala	2890 Gly Phe Ala	Gln ) Asp Pro Gln	Ile Ile Ala Gly 2940 Ser	Arg Ala 2925 Pro	Thr 2910 Ser Pro	2899 Tyr ) Ser Pro	Ser His Val
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 295	Thr Tyr His Lys 2935 Pro	Arg Thr 2920 Thr Gln	Tyr 2905 Gln ) Ala Pro	2890 Gly Phe Ala Val	Gln Asp Pro Gln Gln 2955	Ile Ile Ala Gly 2940 Ser	Arg Ala 2925 Pro Thr	Thr 2910 Ser Pro Gln	2899 Tyr ) Ser Pro	Ser His Val Glu Ala 2960
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2950 Pro Cys	Thr Tyr His Lys 2935 Pro	Arg Thr 2920 Thr Gln	Tyr 2905 Gln ) Ala Pro	2890 Gly Phe Ala Val	Gln Asp Pro Gln Gln 2955	Ile Ile Ala Gly 2940 Ser	Arg Ala 2925 Pro Thr	Thr 2910 Ser Pro Gln	2899 Tyr Ser Pro	Ser His Val Glu Ala 2960 Gln
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2956 Pro Cys 2965	Thr Tyr His Lys 2935 Pro Pro	Thr 2920 Thr Gln Pro	Tyr 2905 Gln Ala Pro	2890 Gly Phe Ala Val Gln 2970	Gln Asp Pro Gln Gln 2955 Leu	Ile Ile Ala Gly 2940 Ser Gly	Arg Ala 2925 Pro Thr	Thr 2910 Ser Pro Gln Pro	Z899 Tyr Ser Pro Pro Gly 2979	Ser  His  Val  Glu  Ala 2960 Gln
Thr Pro Pro Val  Gly Glu Leu Phe 290  Pro Pro Ala Gln 2915  Gly Leu Pro Ser 2930  Gly Glu Pro Leu 2945  Gln Pro Ala Pro  Pro Pro Ser Ser	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2950 Pro Cys 2965 Lys Met	Thr Tyr His Lys 2935 Pro Pro	Thr 2920 Thr Gln Pro	Tyr 2905 Gln Ala Pro Ser	2890 Gly Phe Ala Val Gln 2970 Ser	Gln Asp Pro Gln Gln 2955 Leu	Ile Ile Ala Gly 2940 Ser Gly	Arg Ala 2925 Pro Thr	Thr 2910 Ser Pro Gln Pro	Z895 Tyr Ser Pro Pro Gly 2975 Gly	Ser  His  Val  Glu  Ala 2960 Gln
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 298	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2950 Pro Cys 2965 Lys Met 0	Thr Tyr His Lys 2935 Pro Pro	Arg Thr 2920 Thr Gln Pro Gln	Tyr 2905 Gln Ala Pro Ser Val 2985	2890 Gly Phe Ala Val Gln 2970 Ser	Gln Asp Pro Gln Gln 2955 Leu Gln	Ile Ile Ala Gly 2940 Ser Gly Glu	Arg Ala 2925 Pro Thr Gln Ala	Thr 2910 Ser Pro Gln Pro Lys 2990	Z895 Tyr Ser Pro Pro Gly Z975 Gly	Ser His Val Glu Ala 2960 Gln Thr
Thr Pro Pro Val  Gly Glu Leu Phe 290  Pro Pro Ala Gln 2915  Gly Leu Pro Ser 2930  Gly Glu Pro Leu 2945  Gln Pro Ala Pro  Pro Pro Ser Ser	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2950 Pro Cys 2965 Lys Met 0	Thr Tyr His Lys 2935 Pro Pro	Arg Thr 2920 Thr Gln Pro Gln	Tyr 2905 Gln Ala Pro Ser Val 2985	2890 Gly Phe Ala Val Gln 2970 Ser	Gln Asp Pro Gln Gln 2955 Leu Gln	Ile Ile Ala Gly 2940 Ser Gly Glu	Arg Ala 2925 Pro Thr Gln Ala	Thr 2910 Ser Pro Gln Pro Lys 2990	Z895 Tyr Ser Pro Pro Gly Z975 Gly	Ser His Val Glu Ala 2960 Gln Thr
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro Pro Pro Ser Ser 298	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2950 Pro Cys 2965 Lys Met 0	Thr Tyr His Lys 2935 Pro Pro	Arg Thr 2920 Thr Gln Pro Gln	Tyr 2905 Gln Ala Pro Ser Val 2985 Leu	2890 Gly Phe Ala Val Gln 2970 Ser	Gln Asp Pro Gln Gln 2955 Leu Gln	Ile Ile Ala Gly 2940 Ser Gly Glu	Arg Ala 2925 Pro Thr Gln Ala	Thr 2910 Ser Pro Gln Pro Lys 2990 Ala	Z895 Tyr Ser Pro Pro Gly Z975 Gly	Ser His Val Glu Ala 2960 Gln Thr
Thr Pro Pro Val  Gly Glu Leu Phe	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2956 Pro Cys 2965 Lys Met 0 Glu Gln	Thr Tyr His Lys 2935 Pro Pro	Thr 2920 Thr Gln Pro Gln Arg 3000	Tyr 2905 Gln Ala Pro Ser Val 2985 Leu	2890 Gly Phe Ala Val Gln 2970 Ser	Gln Asp Pro Gln 2955 Leu Gln Gln	Ile Ile Ala Gly 2940 Ser Gly Glu Gly	Arg Ala 2925 Pro Thr Gln Ala Pro 3005	Thr 2910 Ser Pro Gln Pro Lys 2990 Ala	2895 Tyr Ser Pro Pro Gly 2975 Gly	Ser His Val Glu Ala 2960 Gln Thr Arg
Thr Pro Pro Val  Gly Glu Leu Phe	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2956 Pro Cys 2965 Lys Met 0 Glu Gln	Thr Tyr His Lys 2935 Pro Pro Pro	Thr 2920 Thr Gln Pro Gln Arg 3000 Val	Tyr 2905 Gln Ala Pro Ser Val 2985 Leu	2890 Gly Phe Ala Val Gln 2970 Ser	Gln Asp Pro Gln 2955 Leu Gln Gln	Ile Ile Ala Gly 2940 Ser Gly Glu Gly Gln	Arg Ala 2925 Pro Thr Gln Ala Pro 3005 Ala	Thr 2910 Ser Pro Gln Pro Lys 2990 Ala	2895 Tyr Ser Pro Pro Gly 2975 Gly	Ser His Val Glu Ala 2960 Gln Thr Arg
Thr Pro Pro Val  Gly Glu Leu Phe	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2950 Pro Cys 2965 Lys Met 0 Glu Gln His Thr	Thr Tyr His Lys 2935 Pro Pro Pro Gln 3015	Thr 2920 Thr Gln Pro Gln Arg 3000 Val	Tyr 2905 Gln Ala Pro Ser Val 2985 Leu	2890 Gly Phe Ala Val Gln 2970 Ser Pro	Gln Asp Pro Gln 2955 Leu Gln Ala Ala	Ile Ile Ala Gly 2940 Ser Gly Glu Gly Gln 3020	Arg Ala 2925 Pro Thr Gln Ala Pro 3005 Ala	Thr 2910 Ser Pro Gln Pro Lys 2990 Ala Glu	2899 Tyr Ser Pro Pro Gly 2979 Gly Asn	Ser His Val Glu Ala 2960 Gln Thr Arg
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro  Pro Pro Ser Ser 298 Gln Thr Gly Val 2995 Pro Pro Glu Pro 3010 Pro Thr Ser Phe	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 295 Pro Cys 2965 Lys Met 0 Glu Gln His Thr Pro Ser	Thr Tyr His Lys 2935 Pro Pro Pro Gln 3015 Pro	Thr 2920 Thr Gln Pro Gln Arg 3000 Val	Tyr 2905 Gln Ala Pro Ser Val 2985 Leu	2890 Gly Phe Ala Val Gln 2970 Ser Pro	Gln Asp Pro Gln 295! Leu Gln Ala Ala Ser	Ile Ile Ala Gly 2940 Ser Gly Glu Gly Gln 3020 Met	Arg Ala 2925 Pro Thr Gln Ala Pro 3005 Ala	Thr 2910 Ser Pro Gln Pro Lys 2990 Ala Glu	2899 Tyr Ser Pro Pro Gly 2979 Gly Asn	Ser His Val Glu Ala 2960 Gln Thr Arg Gly Leu
Thr Pro Pro Val  Gly Glu Leu Phe	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2956 Pro Cys 2965 Lys Met 0 Glu Gln His Thr Pro Ser 3036	Thr Tyr His Lys 2935 Pro Pro Pro Gln 3015 Pro	Thr 2920 Thr Gln Pro Gln Arg 3000 Val	Tyr 2905 Gln Ala Pro Ser Val 2985 Leu Gln Ser	2890 Gly Phe Ala Val Gln 2970 Ser Pro Arg	Gln Asp Pro Gln 295! Leu Gln Ala Ala Ser 303!	Ile Ile Ala Gly 2940 Ser Gly Glu Gly Gln 3020 Met	Arg Ala 2925 Pro Thr Gln Ala Pro 3005 Ala Lys	Thr 2910 Ser Fro Gln Pro Lys 2990 Ala Glu Pro	2899 Tyr Ser Pro Pro Gly 2979 Gly Asn Thr	Ser His Val Glu Ala 2960 Gln Thr Arg Gly Leu 3040
Thr Pro Pro Val  Gly Glu Leu Phe 290 Pro Pro Ala Gln 2915 Gly Leu Pro Ser 2930 Gly Glu Pro Leu 2945 Gln Pro Ala Pro  Pro Pro Ser Ser 298 Gln Thr Gly Val 2995 Pro Pro Glu Pro 3010 Pro Thr Ser Phe	Val Val 2885 Gln Glu 0 Leu Thr Arg Thr Gln Pro 2955 Pro Cys 2965 Lys Met 0 Glu Gln His Thr Pro Ser 3036 Pro Thr	Thr Tyr His Lys 2935 Pro Pro Pro Gln 3015 Pro	Thr 2920 Thr Gln Pro Gln Arg 3000 Val	Tyr 2905 Gln Ala Pro Ser Val 2985 Leu Gln Ser	2890 Gly Phe Ala Val Gln 2970 Ser Pro Arg Val	Gln Asp Pro Gln 2955 Leu Gln Ala Ala Ser 3035 Lys	Ile Ile Ala Gly 2940 Ser Gly Glu Gly Gln 3020 Met	Arg Ala 2925 Pro Thr Gln Ala Pro 3005 Ala Lys	Thr 2910 Ser Fro Gln Pro Lys 2990 Ala Glu Pro	2899 Tyr Ser Pro Pro Gly 2975 Gly Asn Thr Asp	Ser His Val Glu Ala 2960 Gln Thr Arg Gly Leu 3040 Val
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ttataccaat ataaacaatt actcaggaaa aaaagaaaat aaaaacttgc aagggctaaa

180

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ataacttgct taccaccaaa gatgcttgct ctaagaactg tgaagggatt caagaggaaa
240
agtacaceca gagagggete atacatgtee tetececete etectecace accaggacae
300
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360
tocottotot acttggagag coccatttoa ttacaggaat ttgotttgag ttttattato
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900
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His Thr Glu Thr Ala Ser Ser Phe Gln Pro Ser Pro Phe Ser Ala Asp
                            40
Phe Glu Leu Gln Ile Ser Leu Leu Tyr Leu Glu Ser Pro Ile Ser Leu
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                                            60
Gln Glu Phe Ala Leu Ser Phe Ile Ile Ile Leu Val Tyr Val Leu Asp
                    70
Trp Ala Ala Ile Thr Arg Cys His Arg Leu Ser Gly Leu Asn Asn Lys
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His Ser Tyr Pro Thr Val Thr Glu Ala Glu Lys Pro Gly Val Lys Val
                                105
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Pro Ala Trp Ser Asp Ser Val Leu Glu Ala Gly Lys Ser Lys Met Glu
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                            120
Ala Leu Val Gly Leu Val Ser Gly Arg Ala Ser Leu Cys Phe Gln Asp
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140
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Gly Ala Leu Ser Leu His Leu Pro Glu Gly Arg Asn Ala Val Ser Leu
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Gln His Arg Arg Asn Thr Ser Glu Lys Lys Ser Ser Arg Lys Val Glu
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Asn Lys Glu Met Glu Tyr Ile Tyr Glu Asn Tyr Tyr Ile
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Gly Val Ile Phe Met Asn Gly Asn Arg Ala Ser Glu Ala Val Leu Trp 35 40 45

Glu Ala Leu Arg Lys Met Gly Leu Arg Pro Gly Val Arg His Pro Phe 50 55 60

Leu Gly Asp Leu Arg Lys Leu Ile Thr Asp Asp Phe Val Lys Gln Lys 65 75 80

Tyr Leu Glu Tyr Lys Lys Ile Pro Asn Ser Asn Pro Pro Glu Tyr Glu

85

90

95

Phe Leu Trp Gly Leu Arg Ala Arg His Glu Thr Ser Lys Met Arg Val

Leu Arg Phe Ile Ala Gln Asn Gln Asn Arg Asp Pro Arg Glu Trp Lys
115 120 125

Ala His Phe Leu Glu Ala Val Asp Asp Ala Phe Lys Thr Met Asp Val

Asp Met Ala Glu Glu His Ala Arg Ala Gln Met Arg Ala Gln Met Asn 145 150 155 160

Ile Gly Asp Glu Ala Leu Ile Gly Arg Trp Ser Trp Asp Asp Ile Gln 165 170 175

Val Glu Leu Leu Thr Trp Asp Glu Asp Gly Asp Phe Gly Asp Ala Trp 180 185 190

Ala Arg Ile Pro Phe Ala Phe Trp Ala Arg Tyr His Gln Tyr Ile Leu 195 200 205

Asn Ser Asn Arg Ala Asn Arg Arg Ala Thr Trp Arg Ala Gly Val Ser 210 225 220

Ser Gly Thr Asn Gly Gly Ala Ser Thr Ser Val Leu Asp Gly Pro Ser 225 230 235 240

Thr Ser Ser Thr Ile Arg Thr Arg Asn Ala Ala Arg Ala Gly Ala Ser 245 250 255

Phe Phe Ser Trp Ile Gln 260

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<211> 1731

<212> DNA

<213> Homo sapiens

<400> 4525

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gagacaggga gccaagctag ctcagagcag cctgggcagc taatctcctt cagtgaggcc

ctgcagcact tccagactgt ggacctttcc cccttcaaga aaagaatcca gccaactatt

cgaaggactg ggctcgccgc cctccgacac tacctcttcg ggcctccaaa gctccaccag

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gagtgtetet ccagagagtg taateggeag cagaaggtea teecegtggt gaacagette
720
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cagtgtccca ggtccagaga gcccaaggtg gttgctagac tggttttggc tgcagttctt
ccccatccac actttctcaa attccagctt accaaaatct ccatcaccca ccccctggag
tetgetagtt eteettete tgeeetgaet gtegeeettt tetggtetta taettatgae
aagcatatat totgatcaaa aattgggago cagggtocaa tagttggact attcaaagtt
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1140
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1260
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1380
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1560
gccatgtcag gagcctggcc aggccgcacc ccttgctgtc tcagcagatg ggatatagga
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<211> 344
<212> PRT
<213> Homo sapiens
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<400> 4526

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Glu Ala Val Asp Thr Ile Gln Pro Glu Thr Gly Ser Gln Ala Ser Ser
            40
Glu Gln Pro Gly Gln Leu Ile Ser Phe Ser Glu Ala Leu Gln His Phe
                  55
Gln Thr Val Asp Leu Ser Pro Phe Lys Lys Arg Ile Gln Pro Thr Ile
                   75
              70
Arg Arg Thr Gly Leu Ala Ala Leu Arg His Tyr Leu Phe Gly Pro Pro
                           90 95
            85
Lys Leu His Gln Arg Leu Arg Glu Glu Arg Asp Leu Val Leu Thr Ile
      100 105
Ala Gln Cys Gly Leu Asp Ser Gln Asp Pro Val His Gly Arg Val Leu
     115 120
                                    125
Gln Thr Ile Tyr Lys Lys Leu Thr Gly Ser Lys Phe Asp Cys Ala Leu
                 135
His Gly Asn His Trp Glu Asp Leu Gly Phe Gln Gly Ala Asn Pro Ala
              150
                              155
Thr Asp Leu Arg Gly Ala Gly Phe Leu Ala Leu Leu His Leu Leu Tyr
           165 170
Leu Val Met Asp Ser Lys Thr Leu Pro Met Ala Gln Glu Ile Phe Arg
                                       190
        180 185
Leu Ser Arg His His Ile Gln Gln Phe Pro Phe Cys Leu Met Ser Val
                    200
Asn Ile Thr His Ile Ala Ile Gln Ala Leu Arg Glu Glu Cys Leu Ser
                  215
                                 220
Arg Glu Cys Asn Arg Gln Gln Lys Val Ile Pro Val Val Asn Ser Phe
                              235
              230
Tyr Ala Ala Thr Phe Leu His Leu Ala His Val Trp Arg Thr Gln Arg
           245
                            250
Lys Thr Ile Ser Asp Ser Gly Phe Val Leu Lys Gly Val Leu Phe Leu
        260 265
Leu Gly Arg Pro Arg Leu Asn Ala Gln Cys Pro Arg Ser Arg Glu Pro
      275 280 285
Lys Val Val Ala Arg Leu Val Leu Ala Ala Val Leu Pro His Pro His
   290 295
                                 300
Phe Leu Lys Phe Gln Leu Thr Lys Ile Ser Ile Thr His Pro Leu Glu
305 310 315 320
Ser Ala Ser Ser Pro Phe Ser Ala Leu Thr Val Ala Leu Phe Trp Ser
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Tyr Thr Tyr Asp Lys His Ile Phe
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240
tetteacetg ggaccetegg ceaggetggg acageateea ggaggegagg etgeatggte
cagcggtggg tgcaggtggc aacaggtcgg cgggctgtgc aggttccaaa aggagctctc
gggttggcac tgggtgagac cagccccggg gccagcaggg gaatgagcgg tggagcaggg
gqttqctggg cactggggtg ggccccatct cctgtccttc cctcatggct gctggaaggg
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tgtccagcga ggccatctcc gtggggtcct cagtgttggc gaggaggccg tatcgcctcc
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ageceaqqat cacqtaqaaq gaqegeqtea gegeegagee egaegeeece ggeggaegeg
720
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780
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Ser Gln Lys Gly Ser Leu Gly His Leu Pro Thr Gln Pro Trp Leu Trp
Ala Ala Met Ser Pro Arg Gly Gln Glu Arg Gly Thr Ser His Ser Gln
                                          60
                       55
Ala Arg Glu Pro Gln Arg Pro Gly Arg Trp Leu Leu Gly Ser Leu Gln
                                      75
Ser Ser Pro Gly Thr Leu Gly Gln Ala Gly Thr Ala Ser Arg Arg
                                   90
Gly Cys Met Val Gln Arg Trp Val Gln Val Ala Thr Gly Arg Arg Ala
           100
                               105
Val Gln Val Pro Lys Gly Ala Leu Gly Leu Ala Leu Gly Glu Thr Ser
                                              125
                           120
Pro Gly Ala Ser Arg Gly Met Ser Gly Gly Ala Gly Gly Cys Trp Ala
                       135
Leu Gly Trp Ala Pro Ser Pro Val Leu Pro Ser Trp Leu Leu Glu Gly
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155
                   150
Pro Pro Pro Trp Leu Ser Ile Ile Ser Asp Ser Gly Thr Gln Thr Pro
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               165
Ser Pro Arg Arg Cys Pro Ala Arg Pro Ser Pro Trp Gly Pro Gln Cys
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Trp Arg Gly Gly Arg Ile Ala Ser Ala Glu Ala Ser Ser Thr
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<211> 546
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aagatggagg agaaaccctc agggcccatc ccggacatgc tggccactgc agagcccagc
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240
tecatgtegg aggagecagg ecetgageag geagecacae egecagtggg gaacgtggag
gggctggagg gatgcagcag ggctcctccc cagccccaga cagctgccag tctggccccg
360
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gacgacgccc acctccaggg aagcaaatcc cttgctccag ccctggctgc tgcctcagtt
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540
agtete
546
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Glu Pro Ser Ser Ser Glu Thr Asp Lys Glu Val Leu Ser Pro Ala Val
            20
Pro Ala Ala Pro Ser Ser Ser Met Ser Glu Glu Pro Gly Pro Glu
                            40
Gln Ala Ala Thr Pro Pro Val Gly Asn Val Glu Gly Leu Glu Gly Cys
                        55
Ser Arg Ala Pro Pro Gln Pro Gln Thr Ala Ala Ser Leu Ala Pro Asp
                                        75
                    70
Pro Ala Leu Ala
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<212> PRT
<213> Homo sapiens
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Ser Lys Lys Pro Glu Arg Arg Pro Arg Gly Arg Arg Gly Arg Lys
                       40
Cys Gly Arg Gly His Lys Gly Glu Arg Gln Arg Gly Thr Arg Pro Arg
Leu Gly Phe Glu Gly Gly Gln Thr Pro Phe Tyr Ile Arg Ile Pro Lys
                                 75
                70
Tyr Gly Phe Asn Glu Gly His Ser Phe Arg Arg Gln Tyr Lys Pro Leu
                             90
Ser Leu Asn Arg Leu Gln Tyr Leu Ile Asp Leu Gly Arg Val Asp Pro
                          105
         100
Ser Gln Pro Ile Asp Leu Thr Gln Leu Val Asn Gly Arg Gly Val Thr
                       120
Ile Gln Pro Leu Lys Arg Asp Tyr Gly Val Gln Leu Val Glu Gly
                                    140
                   135
Ala Asp Thr Phe Thr Ala Lys Val Asn Ile Glu Val Gln Leu Ala Ser
                                 155 160
                 150
Glu Leu Ala Ile Ala Ala Ile Glu Lys Asn Gly Gly Val Val Thr Thr
                              170
Ala Phe Tyr Asp Pro Arg Ser Leu Asp Ile Val Cys Lys Pro Val Pro
                           185 190
         180
Phe Phe Leu Arg Gly Gln Pro Ile Pro Lys Arg Met Leu Pro Pro Glu
                       200 205
Glu Leu Val Pro Tyr Tyr Thr Asp Ala Lys Asn Arg Gly Tyr Leu Ala
                    215
                         220
Asp Pro Ala Lys Phe Pro Glu Ala Arg Leu Glu Leu Ala Arg Lys Tyr
225 230 235 240
Gly Tyr Ile Leu Pro Asp Ile Thr Lys Asp Glu Leu Phe Lys Met Leu
             245 250
Cys Thr Arg Lys Asp Pro Arg Gln Ile Phe Phe Gly Leu Ala Pro Gly
          260 265 , 270
Trp Val Val Asn Met Ala Asp Lys Lys Ile Leu Lys Pro Thr Asp Glu
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Asn Leu Leu Lys Tyr Tyr Thr Ser
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<210> 4533
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<212> DNA
<213> Homo sapiens
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Asp Trp Leu Met Gly Lys Ser Lys Ala Lys Pro Asn Gly Lys Lys Pro
                        55
Ala Ala Glu Glu Arg Lys Ala Tyr Leu Glu Pro Glu His Thr Lys Ala
                    70
Arg Ile Thr Asp Phe Gln Phe Lys Glu Leu Val Val Leu Pro Arg Glu
                                     90
Ile Asp Leu Asn Glu Trp Leu Ala Ser Asn Thr Thr Thr Phe Phe His
            100
                                105
His Ile Asn Leu Gln Tyr Ser Thr Ile Ser Glu Phe Cys Thr Gly Glu
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120
       115
Thr Cys Gln Thr Met Ala Val Cys Asn Thr Gln Tyr Tyr Trp Tyr Asp
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                                           140
Glu Arg Gly Lys Lys Val Lys Cys Thr Ala Pro Gln Tyr Val Asp Phe
                   150
                                       155
Val Met Ser Ser Val Gln Lys Leu Val Thr Asp Glu Asp Val Phe Pro
                                   170
Thr Lys Tyr Gly Arg Glu Phe Pro Ser Ser Phe Glu Ser Leu Val Arg
           180
                               185
Lys Ile Cys Arg His Leu Phe His Val Leu Ala His Ile Tyr Trp Ala
                           200
His Phe Lys Glu Thr Leu Ala Leu Glu Leu His Gly His Leu Asn Thr
                       215
                                           220
Leu Tyr Val His Phe Ile Leu Phe Ala Arg Glu Phe Asn Leu Leu Asp
                                      235
                   230
Pro Lys Glu Thr Ala Ile Met Asp Asp Leu Thr Glu Val Leu Cys Ser
               245
                                   250
Gly Ala Gly Gly Val His Ser Gly Gly Ser Gly Asp Gly Ala Gly Ser
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Gly Gly Pro Gly Ala Gln Asn His Val Lys Glu Arg
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Gln Ala Gly Val Gln Trp His Asp His Ser Ser Leu Gln Pro Leu Pro
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Arg Gly Asp Ile Val Phe Phe Leu Gln Lys Val His Ile Pro Glu Ser
Ile Leu Ile Phe Arg Asp Glu Ile Asp Leu His Ala Leu Tyr Gln Ala
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Gly Gln Leu Thr Leu Ile Leu Val Asp His His Ile Leu Ser Lys Ser
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Asp Thr Ala Leu Glu Glu Xaa Ser Ser Arg Gly Ala Arg Pro Ser Thr
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His Arg Ala Glu Thr Leu Pro Ser Leu Xaa His Val Ser Val Glu Leu
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Val Gly Ser Cys Ala Thr Leu Val Thr Glu Arg Ile Leu Gln Gly Ala
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Pro Glu Ile Leu Asp Arg Gln Thr Ala Ala Leu Leu His Gly Thr Ile
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Ile Leu Asp Cys Val Asn Met Asp Leu Lys Ile Gly Lys Ala Thr Pro
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Lys Asp Ser Lys Tyr Val Glu Lys Leu Glu Ala Leu Phe Pro Asp Leu
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Pro Lys Arg Asn Asp Ile Phe Asp Ser Leu Gln Lys Ala Lys Phe Asp
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Val Ser Gly Leu Thr Thr Glu Gln Met Leu Arg Lys Asp Gln Lys Thr
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Lys Leu Gln Gln Glu Gln Arg Gln Val Glu Glu Leu Arg Met Gln Leu
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Gly Tyr Val Ser Leu Gln Glu Lys Asp Ile Phe Val Ser Gly Val Lys
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Ile Thr Ser Asn Arg Leu Gly Arg Ala Pro Val Glu Ser Pro Val Pro
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Ser His Phe Arg Arg Val Ala Leu Leu Pro Arg Ser Arg Ser Gln Trp
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Pro Asp Lys Gln Ser His Ser Gly Val Val Arg Pro Gly Arg Val Ser
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Pro Val Gly Gly Arg Gly Ala Leu Ala Arg Arg Val Ser Gly Glu Ala
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270

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Leu Ile Asp Gly Lys Gly Arg Gly Val Ile Ala Thr Lys Gln Phe Ser
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Arg Gly Asp Phe Val Val Glu Tyr His Gly Asp Leu Ile Glu Ile Thr
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Il $\epsilon$	val	Glu	ı Asr	Gly			His	Pro	Cys			val	TTE	: GIN	Glu
625	5	_		_	630		. mi		, n	635		λ ~~~	- הות	Acn	640 Asn
Ile	Tr	Pro	val			GIU	ıınr	Let	650		nıs	, ALG	что	655	Asn
*	_ <b>_</b> 71-	37-7	C1.	645			2~~	(117			Phe	Ala	Val		Cys
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Gln Glu Arg Asn Leu Glu Glu Lys Ile Lys Gln His Val Leu Gln Met
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Arg Glu Gln Arg Arg Phe His Gly Gln Ala Pro Leu Glu Glu Met Arg
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Lys Ala Ala Glu Asp Leu Glu Ile Ala Thr Glu Leu Gln Asp Glu Val
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Ser Lys Lys Ala Asp Ser Val Ala Ala Lys Val Asp Leu Leu Gly Glu
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Phe Gln Ser Ala Leu Pro Lys Ile Asn Ser His Pro Thr Arg Ser Gln
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Lys Lys Ser Ser Gln Lys Lys Ser Ser Lys Lys Asn His Pro Gln Lys
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Pro Pro Cys His Ile Val Asp Tyr Arg Thr Arg Trp Ser Gly Ile Arg
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Lys Gln His Met Val Asn Ala Thr Pro Phe Lys Ile Ala Arg Gly Gln
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Asn Asp Phe Lys Ala Leu Gln Tyr Phe His Pro Lys Ser Leu Thr Arg
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Gly Phe Arg Pro Gly Met Arg Cys Gly Gly Ser Ser Leu Gly Arg Thr
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Cys Cys Ser Pro Thr Arg Arg Ala Cys Val Val Ser Arg Ala Val Thr
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7	C - 14	7 ~~	180	Cox	The	7) CD	Sor		Len	Ser	Δsn	T.em		Asp	Ser
Asp	ser	195	Ser	Ser	1111	Maii	200	пор	LCu	001		205			
Gl 11	Glu		Leu	Gln	Δla	Lays		Glv	Leu	Lys	Glv		Pro	Glu	His
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T11		Glv	Lvs	T.e11	Glv		Asn	Glv	Glu	Arg		Ala	Glu	Leu	Leu
225	MEC	Gry	цyо	LCu	230			1		235					240
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Leu	GIY	цуз	361.	245	OLY	цуб	<b></b>		250	-1-	<b>-</b> 1	5		255	
ח ז ה	Dro	Lou	Tare		Glv	Gln	Ser	Val	_	Lys	Asp	Val	Ser		Val
Ala	FIQ	шец	260	Val	O± 1	· · · ·		265		-1-	_		270		
Laze	Luc	T. <b>ڪ</b> 11		Gln	Ser	Glv	Glu		Phe	Leu	Gln	Asp	Gly	Ser	Cys
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Ala	Ala	Arg		гуs	rrp	GTY	TIE	505	на	Asn	Cys	PLO	510	110	DCI
•	a1 -	7	500	Cox	17-1	Ι ου	7 ~~		בוג	val	Thr	Acn		Met	Ser
Arg	GIII	515	цур	261	vaı	Беи	520	FIO	ALG	Val		525	1		
<b>71</b> -	T 011		Cor	T10	λen	Dro		בומ	Ser	Ser	Glv		Glu	Thr	Thr
GIII	530		SEL	116	ASII	535	261	AIG	501	501	540	2.0			
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nis	val	-10	пys	565	vah	DCI		rap	570	**** 9				575	
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	Mot	Dro	Thr	Arm		Glu	Asp	ī.eu	Met		Δsn	Len	Pro	Len	
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Glu	Tur	Thr	Lvs		Asp	Glv	Arg	Leu		Leu	Ala	Ser	Ara		Pro
014	- y -	****	900	*** 5		0-1	9	905					910		
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Pro			His	Asp	Pro	Ile	His	Asp	Gln	Ser	Trp	Tyr	Leu	Asp	Gln
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Ser Pro Arg Asn Ser Leu Arg Asn Ile Leu Thr Leu Asn Ser Thr Ala
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Glu Pro Ser Ser Trp Glu Ser Arg Glu Arg Pro Leu Gln Ser Arg Asn
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Val Tyr Ser Ser Ala Ser Phe Ser Glu His Leu Asp Gly Gly Cys Ser
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Lvs	Glu	Val	Glu		Gly	Pro	Gly	Asp	Gln	Gln	Gly	Asp	Ser	Tyr	Leu
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Glu	Lys	Phe	Leu	Arg	His	His		Glu	Thr	Leu	Thr		Ser	Pro	Cys
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	210			_		215	<b>m</b> )	<b>a</b> 1	Dh-	T		602	T 011	Gln.	Luc
	Pro	Arg	Leu	Ser		ser	Thr	GIN	Pne	Leu	Ser	Ser	пеи	GIII	240
225	_	_	D1 .	m1	230	mb so	Dho	Dro	Dro	235 Arg	λ1 =	Thr	Gln	Cvs	
Ala	Ser	Arg	Pne		HIS	1111	Pne	PIU	250	Arg	AIG	1111	0111	255	200
**- 1	T	C 0 26	Dwa	245	v-1	Lvc	T.e.ii	Mot		Arg	Glv	Glv	Ser		Pro
vai	ьуѕ	ser	260	Giu	vai	цуз	цец	265	пор	1119	1	1	270		
7 ~~	715	Glv		Glv	Tyr	Δla	Ser		Asp	Arg	Thr	His	Val	Leu	Ala
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		Gly	Glu	Leu			Leu	GIY	GIN		Leu	GIII	Ala	116	Thr 400
385	_			_	390			<b>a</b>	a1	395	~1 n	~1.,	Dro	Λla	
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Ser	ser			Asp	GIA	ьец	440	ьеи	PIO	FIO	Val	445	****	01	
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 Pro Ser Ala Met Phe Tyr Leu Ala Ala Ala Val Ser Asp Phe Tyr Val
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Gln Lys Lys Pro Val Trp Val Asp Glu Glu Asp Glu Asp Glu Glu Met
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 Val Asp Met Met Asn Asn Arg Phe Arg Lys Asp Met Met Lys Asn Ala
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Thr Ser Thr His Ser Lys Val Leu Tyr Val Tyr Asp Met Leu Ala Gly
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Cys Leu Gln Glu Thr Asn Pro Lys Pro Ile Lys Ala Ile Met Asn Leu
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Ser Leu Asn Ile Ile Tyr Asn Lys Gln Asn Leu Val Asn Leu Gln Lys
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Ser Asn Ala Leu Lys Lys His Gln Ser Leu Cys Met Cys Arg Thr Asp
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Gly Asp Phe Ala Ile Leu Leu Arg Ala Gly Phe Asp Arg Trp Ser Ala
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Asp Ile Val Thr Ile Ser Gln Ala Thr Pro Ser Ser Val Ser Arg Gly
                       55
Thr Ala Pro Ser Asp Asn Arg Val Thr Ser Phe Arg Asp Leu Ile His
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Asp Gln Asp Glu Asp Glu Glu Glu Glu Glu Gly Gln Arg Ser Arg Phe
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                85
Tyr Ala Gly Gly Ser Glu Arg Ser Gly Gln Gln Ile Val Gly Pro Pro
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            100
Arg Lys Lys Ser Pro Asn Glu Leu Val Asp Asp Leu Phe Lys Gly Ala
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        115
Lys Glu His Gly Ala Val Ala Val Glu Arg Val Thr Lys Ser Pro Gly
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                       135
Glu Thr Ser Lys Pro Arg Pro Phe Ala Gly Gly Gly Tyr Arg Leu Gly
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Ala Ala Pro Glu Glu Glu Ser Ala Tyr Val Ala Gly Glu Lys Arg Gln
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                165
His Ser Ser Gln Asp Val His Val Val Leu Lys Leu Trp Lys Ser Gly
                               185
            180
Phe Ser Leu Asp Asn Gly Glu Leu Arg Ser Tyr Gln Asp Pro Ser Asn
                           200
Ala Gln Phe Leu Glu Ser Ile Arg Arg Gly Glu Val Pro Ala Glu Leu
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Arg Arg Leu Ala His Gly Gly Gln Val Asn Leu Asp Met Glu Asp His
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Arg Asp Glu Asp Phe Val Lys Pro Lys Gly Ala Phe Lys Ala Phe Thr
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Gly Glu Gly Gln Lys Leu Gly Ser Thr Ala Pro Gln Val Leu Ser Thr
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265
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Ser Ser Pro Ala Gln Gln Ala Glu Asn Glu Ala Lys Ala Ser Ser Ser
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Ile Leu Ile Asp Glu Ser Glu Pro Thr Thr Asn Ile Gln Ile Arg Leu
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Ala Asp Gly Gly Arg Leu Val Gln Lys Phe Asn His Ser His Arg Ile
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Ser Asp Ile Arg Leu Phe Ile Val Asp Ala Arg Pro Ala Met Ala Ala
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                                    330
Thr Ser Phe Ile Leu Met Thr Thr Phe Pro Asn Lys Glu Leu Ala Asp
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Lys Glu Val Lys Trp Gly Pro Arg Arg Lys Ala Gly Gly Val Trp Ala
Glu Pro Ala Ser Gly Gly Leu Pro Pro Pro Glu Asp Glu Phe Cys Ser
                        55
Pro Gly Val Cys Thr Leu Thr Leu Ala His Ser Leu Thr His Lys Thr
                                        75
Leu Thr Leu Cys Phe Phe Trp Gly Glu Gly Gly His Trp Gln Lys Arg
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Leu Pro Trp Pro Gln Ser Val Pro Ile Leu Ile Phe
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	attttatatt	aaaaggggc	: tcctttttaa	atatatgccg	r tgtaaaaaat
	gaacatctct	: ttgaattgta	tttcttgtat	attacatact	tagagagaga
1800 ctcttttago 1860	: caggcaaagt	cttttttggc	: tgtggctgga	ataaatcatt	: tattacttgg
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395
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385
Glu Asp Gly Gly Ala Leu Arg Gly Glu Val Ile Pro Glu His Glu Phe
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Ala Thr Gly Pro Val Cys Leu Asp Asp Glu Asn Glu Phe Pro Pro Ile
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Ile Leu Cys Arg Gly Asn Gln Lys Gly Lys Thr Lys Gln Ser
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Asp Leu Ser Ser Leu Gln Pro Pro Pro Pro Arg Leu Lys Arg Phe Ser
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His Leu Ser Leu Pro Ser Ser
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 Gln Trp Asn Tyr Cys Thr Leu Ser Gln Glu Ile Leu Arg Arg Pro Ile
                            40
 Val Ala Cys Glu Leu Gly Arg Leu Tyr Asn Lys Asp Ala Val Ile Glu
                                           60
 Phe Leu Leu Asp Lys Ser Ala Glu Lys Ala Leu Gly Lys Ala Ala Ser
                                       75
 His Ile Lys Ser Ile Lys Asn Val Thr Glu Leu Lys Leu Ser Asp Asn
 Pro Ala Trp Glu Gly Asp Lys Gly Asn Thr Lys Gly Asp Lys His Asp
                                105
            100
 Asp Leu Gln Arg Ala Arg Phe Ile Cys Pro Val Val Gly Leu Glu Met
                                               125
                            120
 Asn Gly Arg His Arg Phe Cys Phe Leu Arg Cys Cys Gly Cys Val Phe
                        1.35
 Ser Glu Arg Ala Leu Lys Glu Ile Lys Ala Glu Val Cys His Thr Cys
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160

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155
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145
Gly Ala Ala Phe Gln Glu Asp Asp Val Ile Met Leu Asn Gly Thr Lys
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                                    170
Glu Asp Val Asp Val Leu Lys Thr Arg Met Glu Glu Arg Arg Leu Arg
                                185
            180
Ala Lys Leu Glu Lys Lys Thr Lys Lys Pro Lys Ala Ala Glu Ser Val
                                                205
        195
                            200
Ser Lys Pro Asp Val Ser Glu Glu Ala Pro Gly Pro Ser Lys Val Lys
                        215
Thr Gly Lys Pro Glu Glu Ala Ser Leu Asp Ser Arg Glu Lys Lys Thr
                                        235
                    230
Asn Leu Ala Pro Lys Ser Thr Ala Met Asn Glu Ser Ser Ser Gly Lys
                                    250
                245
Ala Gly Lys Pro Pro Cys Gly Ala Thr Lys Arg Ser Ile Ala Asp Ser
                                265
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Lys Arg Ser Lys Glu Glu Ser Ala His Trp Val Thr His Thr Ser Tyr
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Cys Phe
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720
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780
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<213> Homo sapiens
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Leu Glu Gln Glu Leu Pro Gly Ala Val Phe Ile Leu Cys Asp Val Thr
                      55
Gln Glu Asp Asp Met Lys Thr Leu Val Ser Glu Thr Ile Arg Arg Phe
                                      75
Gly Arg Leu Asp Cys Val Val Asn Asn Ala Gly His His Pro Pro Pro
                                  90
              85
Gln Arg Pro Glu Glu Thr Ser Ala Gln Gly Phe Arg Gln Leu Leu Glu
                              105
Leu Asn Leu Leu Gly Thr Tyr Thr Leu Thr Lys Leu Ala Leu Pro Tyr
                          120
Leu Arg Lys Ser Gln Gly Asn Val Ile Asn Ile Ser Ser Leu Val Gly
                      135
                                         140
Ala Ile Gly Gln Ala Gln Ala Val Pro Tyr Val Ala Thr Lys Gly Ala
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Val Thr Ala Met Thr Lys Ala Leu Ala Leu Asp Glu Ser Pro Tyr Gly
                                  170
Val Arq Val Asn Cys Ile Ser Pro Gly Asn Ile Trp Thr Pro Leu Trp
                              185
Glu Glu Leu Ala Ala Leu Met Pro Asp Pro Arg Ala Thr Ile Arg Glu
                           200
Gly Met Leu Ala Gln Pro Leu Gly Arg Met Gly Gln Pro Ala Glu Val
                       215
                                          220
Gly Ala Ala Ala Val Phe Leu Ala Ser Glu Ala Asn Phe Cys Thr Gly
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                                      235
Ile Glu Leu Leu Val Thr Gly Gly Ala Glu Leu Gly Tyr Gly Cys Lys
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	cctcacaggg	ccgggtgggc	tggcgagccg	acgcggcggc	ggaggaggct
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Ala	Arg	Lys	Tyr	Thr	Ile	Pro		Asp	Hıs	ile	GTĀ	Pne	. GIU	. Phe	Glu
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cctcgtgcac gtgctgcagc tgaagaaccc ggcggggctg gcggtgaagg aagactgcaa
agtccacatc cgagtctatt tgcccccact tcggtggata gcggctgtag caactgcacc
360
cagaccagec etecgtacec agagecetgt tgcatgggta tegactecat eetgggecae
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gttgataagg aaaccaacac ggaagatctc tttctggaag aagcagccag cctcgtgaag
gageggeeca geegeeggge eegagggteg cettttgtte ggagtggeae gattgteegt
teccagaeat tetegeetgg ageaegaage cagtatgttt geagaettta tegtagtgae
660
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agegacagtt caacgetgee ceggaagtee ceetttgtee gaaataettt ggaaagaega
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780
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Pro Tyr Ser Pro Glu Lys Phe Gln Pro Ser Pro Leu Lys Val Asp Lys
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Glu Thr Asn Thr Glu Asp Leu Phe Leu Glu Glu Ala Ala Ser Leu Val
Lys Glu Arg Pro Ser Arg Arg Ala Arg Gly Ser Pro Phe Val Arg Ser
                        55
Gly Thr Ile Val Arg Ser Gln Thr Phe Ser Pro Gly Ala Arg Ser Gln
65
                    70
Tyr Val Cys Arg Leu Tyr Arg Ser Asp Ser Asp Ser Ser Thr Leu Pro
                                    90
                85
Arg Lys Ser Pro Phe Val Arg Asn Thr Leu Glu Arg Arg Thr Leu Arg
                                105
Tyr Lys Gln Ser Cys Arg Ser Ser Leu Ala Glu Leu Met Ala Arg Thr
                            120
Ser Leu Asp Leu Glu Leu Asp Leu Gln Ala Ser Arg Thr Arg Gln Arg
                        135
                                            140
Gln Leu Asn Glu Glu Leu Cys Ala Leu Arg Glu Leu Arg Gln Arg Leu
                    150
                                        155
Glu Asp Ala Gln Leu Arg Gly Gln Thr Asp Leu Pro Pro Trp Val Leu
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Arg Asp Glu Arg Leu Arg Gly Leu Leu Arg Glu Ala Glu Arg Gln Thr
                                185
Arg Gln Thr Lys Leu Asp Tyr Arg His Glu Gln Ala Ala Glu Lys Met
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200

195

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Leu Lys Lys Ala Ser Lys Glu Ile Tyr Gln Leu Arg Gly Gln Ser His
                                          220
                      215
Lys Glu Pro Ile Gln Val Gln Thr Phe Arg Glu Lys Ile Ala Phe Phe
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Thr Arg Pro Arg Ile Asn Ile Pro Pro Leu Pro Ala Asp Asp Val
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cttgatctcc agcacgaaga tgtaaaggaa ccacaggatc atggcgtagc cgcgcttggc
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gtegeceace ageacgatga tgeacaegee gatettgege gggecetggt tetgeteeae
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Gln Gln Gln Arg Gln Arg Leu Ala Arg His Gly Val Arg Arg Ala Ala
Pro Arg Arg Leu Val Val Leu Glu Asp Glu Val Glu Leu Asp Leu Gln
His Glu Asp Val Lys Glu Pro Gln Asp His Gly Val Ala Ala Leu Gly
                                       75
                   70
Arg Ala His Leu Gly Ala His Pro His Gly His Val Ala Gln His Gln
                                   90
Gln Glu Ala His Val Ala His Gln His Asp Asp Ala His Ala Asp Leu
            100
                               105
Ala Arg Ala Leu Val Leu Leu His Gln Val Arg Val His Asp Gly His
                                               125
                           120
Ala Ala His Asp His Gln Arg Gly Gln Ala His Val Ala Pro Val Arg
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140
Gly Arg Gln His His Gly Arg Pro
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145
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<212> DNA
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gagtcaggcc tagggaaatc caccctcatc aacagcctct tcctcaccaa cctctatgag
gategecagg tgecagagge cagtgetege ttgacacaga ccctggccat tgagegeegg
ggcgtagaga ttgaggaagg gggtgtgaaa gtgaagctga cccttgtgga cacacctggc
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tttggggact cagtggactg ctctgactgc tggcttccgg tggtgaaatt catcgaggag
caatttgagc agtaccttag ggatgagagt ggcctgaacc ggaagaacat ccaggactcc
420
egagtecact getgeeteta etteatetea eeetteggee gggeteegge eeetagatgt
480
qqcttcctcc qqqcaataca cgagaaagtc aacatcatcc cagtcattgg caaagcggat
gecetgatge eccaggaaac ecaggeeete aageagaaga teegggatea gttgaaggaa
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720
gta
723
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Arg Lys Ser Val Lys Lys Gly Phe Asp Phe Thr Leu Met Val Ala Gly
                                25
Glu Ser Gly Leu Gly Lys Ser Thr Leu Ile Asn Ser Leu Phe Leu Thr
                            40
Asn Leu Tyr Glu Asp Arg Gln Val Pro Glu Ala Ser Ala Arg Leu Thr
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Gln Thr Leu Ala Ile Glu Arg Arg Gly Val Glu Ile Glu Gly Gly
                    70
                                        75
Val Lys Val Lys Leu Thr Leu Val Asp Thr Pro Gly Phe Gly Asp Ser
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95
                                    90
Val Asp Cys Ser Asp Cys Trp Leu Pro Val Val Lys Phe Ile Glu Glu
                                105
            100
Gln Phe Glu Gln Tyr Leu Arg Asp Glu Ser Gly Leu Asn Arg Lys Asn
                            120
Ile Gln Asp Ser Arg Val His Cys Cys Leu Tyr Phe Ile Ser Pro Phe
                                            140
                        135
Gly Arg Ala Pro Ala Pro Arg Cys Gly Phe Leu Arg Ala Ile His Glu
                                        155
Lys Val Asn Ile Ile Pro Val Ile Gly Lys Ala Asp Ala Leu Met Pro
                                    170
                165
Gln Glu Thr Gln Ala Leu Lys Gln Lys Ile Arg Asp Gln Leu Lys Glu
                                185
Glu Glu Ile His Ile Tyr Gln Phe Pro Glu Cys Asp Ser Asp Glu Asp
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Glu Asp Phe Lys Arg Gln Asp Ala Glu Met Lys Glu Ser Ile Pro Phe
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Ala Val Val Gly Ser Cys Glu Val Val
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<212> DNA
<213> Homo sapiens
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120
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caactggcag acaggcatgt gtgactgttt cagcgactgc ggagtctgtc tctgtggcac
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 tataaaccaa atgaaatatt ttactgataa gattcttcat gcttctttgc tctccttaaa
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864
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<211> 192
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Met Pro Ser Val Val Leu Lys His Ile His His Ile Ser Val Ala Lys
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Asp Gly Glu Glu Leu Lys Leu Lys Arg Cys Leu Leu Asn Phe Val Ala
                               25
Ser Val Arg Ala Phe His His Gln Phe Leu Glu Ser Thr His Gly Ser
                           40
Pro Ser Val Asp Ile Ser Leu Asp Leu Ala Lys Ser Thr Met Arg Thr
                       55
Ala Lys Ser Cys His Ile Val Ile Thr Asn Arg Ser Arg Asp Ala Ile
                   70
Ser Gly Pro Val Glu Ser Pro His Cys Asp Ala Cys Ser Thr Gln Thr
               85
                                   90
Ala Phe Ile His Ile Ser Cys Asn Leu Thr Pro Lys Ala Arg Glu Thr
                               105
           100
Lys Cys Ala Thr Glu Thr Asp Ser Ala Val Ala Glu Thr Val Thr His
                           120
Ala Cys Leu Pro Val Gly Val Leu Gly Gly Arg Thr Gly Thr Asp Ser
                                            140
                       135
Arg Leu Gly His Asn Asp His Arg Arg Leu Ser Leu His Phe Gln Cys
                                       155
                   150
Arg Ala Phe His Val Val Phe Ile Cys Gly Glu Ile Leu Ser Gln Ala
                                   170
Thr Arg His Phe Leu Leu Gly Thr Leu Phe Thr Asn Phe His Cys Phe
                              185
           180
<210> 4661
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<212> DNA
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153
<210> 4662
<211> 51
<212> PRT
<213> Homo sapiens
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Tyr Met Ile Ser Lys His Ser His Glu Gln Ser Asp Arg Gly Glu Gly
                                25
Val Glu Val Val Gln Asn Glu Pro Phe Glu Asp Pro His His Gly His
                                                45
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                            40
Gly Gln Phe
    50
<210> 4663
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<212> DNA
<213> Homo sapiens
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cagacggatg acccaggece ectegatgge ectgacetee aggecageca etcagagete
caggtgccca cccctggcag agccggccta ctgaacacct ctggtaccaa aggcttagaa
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420
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tccacagage tettecaget aaggaaceag etgaaggaee tgaagaeeet ggagageeag
aacctgttct gctgcctgta ccgctcctgg tgccacaacc cagtcaccac ggtgtccctc
600
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cggctccagt gcgtgcccaa ccctgagctg ctgcagaccg aagacagtct aaaggcagcc
cccaagtccc agaaagctga ctcccctagc atcgactacg cagagctgct gcagcacttt
960
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catgaaacac taagggtcgt cacgccctcc cgaggagetc aaggacetge ctgtcaggac
cagggctggg cctgccaacc cagggcagtg ttggggccgg aggctgctgt gtctgcccaa
1200
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getectetea gagtecagte eccaggeete cagegetgte agetgeacce tggeattete acaqaqetqq etqeecacce agtqggggge tatageetca gagaccacte atectetgga 1320 atcaacctct ttctaatacc ctcttggaaa aagagcttgc ccctcctcca gcacactaga qctctqqcct tqtqtqtata tgtatacata cgtgaacaca tgcctgtgtg tgtgtgtgt tgtgtacttg tatgcacgta ggcaccagca caaagatctg aatgatgcac cccacccca 1550 <210> 4664 <211> 347 <212> PRT <213> Homo sapiens <400> 4664 Met Phe Arg His Thr Asp Ser Leu Phe Pro Ile Leu Leu Gln Thr Leu 10 Ser Asp Glu Ser Asp Glu Val Ile Leu Lys Asp Leu Glu Val Leu Ala Glu Ile Ala Ser Ser Pro Ala Gly Gln Thr Asp Asp Pro Gly Pro Leu Asp Gly Pro Asp Leu Gln Ala Ser His Ser Glu Leu Gln Val Pro Thr 55 Pro Gly Arg Ala Gly Leu Leu Asn Thr Ser Gly Thr Lys Gly Leu Glu 75 70 Cys Ser Pro Ser Thr Pro Thr Met Asn Ser Tyr Phe Tyr Lys Phe Met 90 Ile Asn Leu Leu Lys Arg Phe Ser Ser Glu Arg Lys Leu Leu Glu Val 105 100 Arg Gly Pro Phe Ile Ile Arg Gln Leu Cys Leu Leu Leu Asn Ala Glu 120 Asn Ile Phe His Ser Met Ala Asp Ile Leu Leu Arg Glu Glu Asp Leu 135 140 Lys Phe Ala Ser Thr Met Val His Ala Leu Asn Thr Ile Leu Leu Thr 150 155 Ser Thr Glu Leu Phe Gln Leu Arg Asn Gln Leu Lys Asp Leu Lys Thr 170 Leu Glu Ser Gln Asn Leu Phe Cys Cys Leu Tyr Arg Ser Trp Cys His 190 185 Asn Pro Val Thr Thr Val Ser Leu Cys Phe Leu Thr Gln Asn Tyr Arg 200 His Ala Tyr Asp Leu Ile Gln Lys Phe Gly Asp Leu Glu Val Thr Val 220 215 Asp Phe Leu Ala Glu Val Asp Lys Leu Val Gln Leu Ile Glu Cys Pro 230 235 Ile Phe Thr Tyr Leu Arg Leu Gln Leu Leu Asp Val Lys Asn Asn Pro 245 250 Tyr Leu Ile Lys Ala Leu Tyr Gly Leu Leu Met Leu Leu Pro Gln Ser 265 Ser Ala Phe Gln Leu Leu Ser His Arg Leu Gln Cys Val Pro Asn Pro

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275
                           280
Glu Leu Leu Gln Thr Glu Asp Ser Leu Lys Ala Ala Pro Lys Ser Gln
                       295
Lys Ala Asp Ser Pro Ser Ile Asp Tyr Ala Glu Leu Leu Gln His Phe
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Glu Lys Val Gln Asn Lys His Leu Glu Val Arg His Gln Arg Ser Gly
                                  330
               325
Arg Gly Asp His Leu Asp Arg Arg Val Val Leu
                               345
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<211> 1043
<212> DNA
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tgtccacctt tacgaagccg agcatacaca ccacctgaag atctccagag tcgtttggaa
tettacgtta aagaagtttt tggtteatet etteetagta attggeaaga eateteeetg
300
gaagatagtc gtctaaagtt caatcttctg gctcatttag ctgatgactt gggtcatgta
360
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gtccctattc aagatagatc taaatttgat gaactcagtg ccagtaatct gccccccaat
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cagaactgtt ctctaaaccc actttttctg tagaggaatg tatcatcttt ttttttctca
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<211> 167

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                                25
Arg Glu Phe Trp Ser Arg Phe Arg Lys Glu Lys Glu Pro Val Val Val
Glu Thr Val Glu Glu Lys Lys Glu Pro Ile Leu Val Cys Pro Pro Leu
                        55
Arg Ser Arg Ala Tyr Thr Pro Pro Glu Asp Leu Gln Ser Arg Leu Glu
                    70
                                        75
Ser Tyr Val Lys Glu Val Phe Gly Ser Ser Leu Pro Ser Asn Trp Gln
                85
                                    90
Asp Ile Ser Leu Glu Asp Ser Arg Leu Lys Phe Asn Leu Leu Ala His
                                105
            100
Leu Ala Asp Asp Leu Gly His Val Val Pro Asn Ser Arg Leu His Gln
                            120
Met Cys Arg Val Arg Asp Val Leu Asp Phe Tyr Asn Val Pro Ile Gln
                        135
Asp Arg Ser Lys Phe Asp Glu Leu Ser Ala Ser Asn Leu Pro Pro Asn
                                        155
                    150
Leu Lys Ile Thr Trp Ser Tyr
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<210> 4667
<211> 1031
<212> DNA
<213> Homo sapiens
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420
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tgcatgaggc tgatcgggtt ctcagagagg gccctggcac tcatgaaggc ccgcgtgagt
600
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gettteecce geacceagea etgacteaga accaceacet tetgetttge tgteggaett
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aagaagttgc attcctgtct gctttgcatc tgctactttg ctgcagtttg gattcagagc
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840
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aaaaaaaaa a
1031
<210> 4668
<211> 207
<212> PRT
<213> Homo sapiens
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Ala Pro Asp Thr Gly Asn Met Glu Leu Leu Val Arg Tyr Gly Thr Glu
Ala Gln Lys Ala Arg Trp Leu Ile Pro Leu Leu Glu Gly Lys Ala Arg
                             40
Ser Cys Phe Ala Met Thr Glu Pro Gln Val Ala Ser Ser Asp Ala Thr
                         55
Asn Ile Glu Ala Ser Ile Arg Glu Glu Asp Ser Phe Tyr Val Ile Asn
                                         75
                     70
Gly His Lys Trp Trp Ile Thr Gly Ile Leu Asp Pro Arg Cys Gln Leu
                                     90
Cys Val Phe Met Gly Lys Thr Asp Pro His Ala Pro Arg His Arg Gln
                                                     110
                                 105
             100
 Gln Ser Val Leu Leu Val Pro Met Asp Thr Pro Gly Ile Lys Ile Ile
                                                 125
                             120
 Arg Pro Leu Thr Val Tyr Gly Leu Glu Asp Ala Pro Gly Gly His Gly
                                             140
                         135
 Glu Val Arg Phe Glu His Val Arg Val Pro Lys Glu Asn Met Val Leu
                                         155
 Gly Pro Gly Arg Gly Phe Glu Ile Ala Gln Gly Arg Leu Gly Pro Gly
                                     170
                 165
 Arg Ile His His Cys Met Arg Leu Ile Gly Phe Ser Glu Arg Ala Leu
                                 185
 Ala Leu Met Lys Ala Arg Val Ser Ala Phe Pro Arg Thr Gln His
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 <210> 4669
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 <212> DNA
 <213> Homo sapiens
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600
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gattcaagag tggtataaag ctt
683
<210> 4670
<211> 135
<212> PRT
<213> Homo sapiens
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Xaa Ser Phe Ser Gly Leu Arg Gly Ile Ile Gln Glu Lys Tyr Arg Ala
                                     10
Asn Lys Lys Gln Lys Val Phe Gln His Asn Glu Leu Lys Lys Glu
                                 25
Thr Cys Val Gln Ala Gly Phe Gln Asp Met Asn Ile Lys Lys Gln Ile
Gln Glu Gln His Gln Ala Ala Ile Ile Ile Gln Lys His Cys Lys Ala
                                             60
 Phe Lys Ile Arg Lys His Tyr Leu His Ile Arg Ala Thr Val Val Ser
                                         75
 Ile Gln Arg Arg Tyr Arg Lys Leu Thr Ala Val Arg Thr Gln Ala Val
                 85
 Ile Cys Ile Gln Ser Tyr Tyr Arg Gly Phe Lys Val Arg Lys Asp Ile
                                 105
 Gln Asn Met His Arg Ala Ala Thr Leu Ile Gln Ser Phe Tyr Arg Met
                             120
 His Arg Ala Lys Val Asp Tyr
                         135
     130
 <210> 4671
 <211> 657
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<212> DNA
<213> Homo sapiens
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Arg Gly Ser Leu Ser Ser Ser Ser Ser Ser Ser Ser Leu Thr Lys
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Ser Pro Cys Ser Leu Thr Phe Ser Arg Ala Ile Lys Ala Thr Ser Ser
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 Gly Pro Val Ser Gly His Leu Val Ile
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315

305

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 Ser His Tyr Glu Phe Ser Arg Val Arg Glu Phe Val Gly Gln Leu Val
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 Gly Glu Ala Ala Gln Asp Ala Val Arg Ala Ser Ala Gln Arg Met Gly
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Arg Gly Ser Ile Leu Asp Ala Met Arg Pro Gln Gln Leu His Ala Thr
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Glu Ile Thr Ser Ser Gly Phe Arg Leu Ala Trp Pro Pro Leu Leu Thr
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Val Leu Gly Phe Ala Val Gly Thr Cys Thr Gly Ile Tyr Ala Ala Gln
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100

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Val Ala Ala Val Ser Ile Thr Gly Arg Lys Arg Ser Arg Val Ala Pro
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Pro Arg Ser Xaa Met Ala Leu Val Leu Glu Arg Val Cys Ser Thr Leu
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Leu Gly Leu Glu Glu His Leu Asn Ala Leu Asp Arg Ala Ala Gly Asp
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Gly Asp Cys Gly Thr Thr His Ser Arg Ala Ala Arg Ala Ile Gln Glu
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Ala Asp Leu Leu Gln Val Leu Thr Lys Ala Val Lys Ser Ala Glu Ala
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Ala Ala Glu Ala Thr Lys Asn Met Glu Ala Gly Ala Gly Arg Ala Ser
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Tyr Ile Ser Ser Ala Arg Leu Glu Gln Pro Asp Pro Gly Ala Val Ala
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Ala Gln Cys Val Thr Val Lys Glu Lys Leu Leu Glu Gln Ala Glu Ser
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Leu Leu Ser Glu Pro Met Val His Glu Leu Val Leu Trp Ile Gln Gln
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Lys Cys Thr Phe Ser Thr Ser Thr Thr Met Asp Asp Gly Leu Trp Ile
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Tyr Phe Gly Ala Gln Ser Val Arg Val Leu Ser Asp Lys Gly Arg Leu
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Lys His Thr Ala Phe Ala Thr Phe Pro Asn Glu Lys Ala Ala Ile Lys
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Tyr Phe Gly Ala Gln Ser Val Arg Val Leu Ser Asp Lys Gly Arg Leu
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Lys His Thr Ala Phe Ala Thr Phe Pro Asn Glu Lys Ala Ala Ile Lys
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Gln Ser Arg Gly Phe Gly Phe Val Lys Phe Lys Asp Pro Asn Cys Val
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Gly Thr Val Leu Ala Ser Arg Pro His Thr Leu Asp Gly Arg Asn Ile
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Pro Lys Glu Gly Trp Gln Lys Gly Pro Arg Ser Asp Asn Ser Lys Ser
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Asn Lys Ile Phe Val Gly Gly Ile Pro His Asn Cys Gly Glu Thr Glu
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40

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Arg Glu Glu Glu Glu Asn Asp Asp Asn Ser Leu Glu Gly Glu
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Pro Gln Thr Asp Arg Leu Thr Cys Pro Lys Gly Leu Pro Trp Ala Pro
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Lys Val Arq Glu Lys Asp Ile Glu Met Phe Leu Glu Ser Ser Arg Ser
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Lys Phe Ile Gly Tyr Thr Leu Gly Ser Asp Thr Asn Thr Val Val Gly
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Tyr Thr Ser Ile Ala Glu Val Gln Ala Gln Met Lys Glu Glu Tyr Leu
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Arg Ser Pro Leu Ser Gly Gly Glu Glu Glu Val Glu Gln Val Pro Ala
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Thr Asp Ser Ile Asn Ile Leu Ala Asp Val Leu Pro Glu Glu Met Pro
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Gln Gly Ala Pro Gly Phe Pro Lys Asp Met Asp Leu Ala Cys Lys Tyr
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Leu Thr Gly Glu Ser Glu Ser Ser Ser Glu Asp Glu Phe Glu Lys Glu
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Met Glu Ala Glu Leu Asn Ser Thr Met Lys Thr Met Glu Asp Lys Leu
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Asp Glu Asp Arg Ala Val Gln Val Thr Lys Lys Lys Lys Lys Gln
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Asn Arg Lys Lys Arg Arg Val His Lys Lys Met Arg Ser Asn Arg Glu
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Val Ala Gly Ala His Gly Leu Leu Cys Leu Leu Ser Asp His Val Asp
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Lys Arg Ile Leu Asp Ala Ala Gly Ala Asn Leu Lys Val Ile Ser Thr
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Met Ser Val Gly Ile Asp His Leu Ala Leu Asp Glu Ile Lys Lys Arg
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 Gly Ile Arg Val Gly Tyr Thr Pro Asp Val Leu Thr Asp Thr Thr Ala
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             100
 Glu Leu Ala Val Ser Leu Leu Leu Thr Thr Cys Arg Arg Leu Pro Glu
                                                 125
                             120
 Ala Ile Glu Glu Val Lys Asn Gly Gly Trp Thr Ser Trp Lys Pro Leu
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 Trp Leu Cys Gly Tyr Gly Leu Thr Gln Ser Thr Val Gly Ile Ile Gly
                                         155
                     150
 Leu Gly Arg Ile Gly Gln Ala Ile Ala Arg Arg Leu Lys Pro Phe Gly
                                     170
                 165
 Val Gln Arg Phe Leu Tyr Thr Gly Arg Gln Pro Arg Pro Glu Glu Ala
                                                     190
                                 185
 Ala Glu Phe Gln Ala Glu Phe Val Ser Thr Pro Glu Leu Ala Ala Gln
                                                 205
                             200
 Ser Asp Phe Ile Val Val Ala Cys Ser Leu Thr Pro Ala Thr Glu Gly
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220
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Leu Cys Asn Lys Asp Phe Phe Gln Lys Met Lys Glu Thr Ala Val Phe
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Ile Asn Ile Ser Arg Gly Asp Val Val Asn Gln Asp Asp Leu Tyr Gln
                                    250
Ala Leu Ala Ser Gly Lys Ile Ala Ala Ala Gly Leu Asp Val Thr Ser
                                                     270
                                265
            260
Pro Glu Pro Leu Pro Thr Asn His Pro Leu Leu Thr Leu Lys Asn Cys
                            280
Val Ile Leu Pro His Ile Gly Ser Ala Thr His Arg Thr Arg Asn Thr
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                                             300
Met Ser Leu Leu Ala Ala Asn Asn Leu Leu Ala Gly Leu Arg Gly Glu
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Pro Met Pro Ser Glu Leu Lys Leu
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Lys Gln Ala Ala Leu Lys Ser His Tyr Ala Asp Val Asp Pro Glu Asn
Gln Asn Phe Leu Leu Glu Ser Asn Leu Gly Lys Lys Lys Tyr Glu Thr
Glu Phe His Pro Gly Thr Thr Ser Phe Gly Met Ser Val Phe Asn Leu
                                        75
                    70
Ser Asn Ala Ile Val Gly Ser Gly Ile Leu Gly Leu Ser Tyr Ala Met
Ala Asn Thr Gly Ile Ala Leu Phe Ile Ile Leu Leu Thr Phe Val Ser
                                105
            100
Ile Phe Ser Leu Tyr Ser Val His Leu Leu Leu Lys Thr Ala Asn Glu
                            120
Gly Gly Ser Leu Leu Tyr Glu Gln Leu Gly Tyr Lys Ala Ser Gly Leu
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Val Gly Lys Leu
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<212> DNA
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Ala Arg Met Ala Gly His Val Ser Val Leu Val Ser His Phe Pro Pro
                            40
                                                 45
Ser Val Thr Tyr Leu Gly Ile Pro Gln Gly Leu Leu Glu Cys Asp Cys
                                            60
                        55
Pro Leu Pro Ser Cys Leu Gly Tyr Lys Ser Trp Pro Tyr Val Pro Ala
                                        75
Val Arg Gly Ser Gly Asn Pro Thr Gln Pro Pro Val Leu Gly Trp Ser
                                    90
                85
Val Ser Ile His Pro Leu Val Val Ile Glu Ala Ala Leu Pro Val Leu
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Gly Glu Asp Ile Trp Ala Thr Arg Ala Pro Leu Ala Pro Ser Arg Arg
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gcatgcggtg gtgaagatgg attactgaaa gttttgaaat tagagacgca gacagatgat
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<213> Homo sapiens
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Val Glu Gly Leu Ser Gly Arg Arg Asp Pro Leu Gly Asp Pro Thr Met
Phe Phe Tyr Leu Ser Lys Lys Ile Ser Ile Pro Asn Asn Val Lys Leu
                           40
Gln Cys Val Ser Trp Asn Lys Glu Gln Gly Phe Ile Ala Cys Gly Gly
                                            60
                       55
Glu Asp Gly Leu Leu Lys Val Leu Lys Leu Glu Thr Gln Thr Asp Asp
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                   70
Ala Lys Leu Arg Gly Leu Ala Ala Pro Ser Asn Leu Ser Met Asn Gln
                                   90
                85
Thr Leu Glu Gly His Ser Gly Ser Val Gln Val Val Thr Trp Asn Glu
                                105
            100
Gln Tyr Gln Lys Leu Thr Thr Ser Asp Glu Asn Gly Leu Ile Ile Val
                            120
Trp Met Leu Tyr Lys Gly Ser Trp Ile Glu Glu Met Ile Asn Asn Arg
                                            140
                       135
Asn Lys Ser Val Val Arg Ser Met Ser Trp Asn Ala Asp Gly Gln Lys
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Ile Cys Ile Val Tyr Glu Asp Gly Ala Val Ile Val Gly Ser Val Asp
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Gly Asn Arg Ile Trp
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 <212> PRT
 <213> Homo sapiens
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Glu Asn Pro Glu Trp Glu Lys Ala Arg Gln Ala Leu Ala Ser Ile Ser
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Lys Ser Gly Ala Ala Gly Gly Ser Ala Lys Ser Ser Ser Asn Gly Pro
                            40
Val Ala Ser Ala Gln Tyr Val Ser Gln Ala Lys Ala Ser Ala Leu Gln
Gln Gln Gln Tyr Tyr Gln Trp Tyr Gln Gln Asp Asn Tyr Ala Tyr Pro
Tyr Ser Tyr Tyr Pro Met Pro Pro Gly Pro Gly Met
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<212> DNA
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            20
Pro Glu Gly Gly Val Ser Lys Phe Ser Pro Pro Lys Asn Gln Ile Leu
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 Ser Phe Ile Pro Pro Pro Phe Pro Pro Phe Gly Phe Phe Lys Lys Phe
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Pro Ser Phe Phe Arg Lys Gly Lys Gly Glu Arg Gly Gly Gln Arg
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Lys Thr Pro Phe Phe Leu Arg Lys Lys Arg Glu Lys Lys Lys
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 Lys Glu Arg Lys Thr Pro Val Asp Leu Arg Glu Val Asn
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Lys Ala Pro Ala Gly Asp Gly Ser Gln Thr Arg Gly Lys Met Ser Glu
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Gly Gly Arg Lys Ser Ser Leu Leu Gln Lys Ser Lys Ala Asp Ser Ser
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Gly Val Gly Lys Gly Asp Leu Gln Ser Thr Leu Leu Glu Gly His Gly
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Thr Ala Pro Pro Asp Leu Asp Leu Ser Ala Ile Asn Asp Lys Ser Ile
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                                 105
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Val Lys Lys Thr Pro Gln Leu Ala Lys Thr Ile Ser Lys Lys Pro Glu
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Ser Ala Gly Ile Gln Arg Ala Gln Ile Gln Lys Glu Leu Trp Arg Ile
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Gln Asp Val Met Glu Gly Leu Ser Lys His Lys Gln Gln Arg Gly Thr
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Thr Glu Ile Gly Met Ile Gly Ser Lys Pro Phe Ser Thr Val Lys Tyr
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Lys Asn Glu Gly Pro Asp Tyr Arg Leu Tyr Lys Ser Glu Pro Glu Leu
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Thr Thr Val Ala Glu Val Asp Glu Ser Asn Gly Glu Glu Lys Ser Glu
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Pro Val Ser Glu Ile Glu Thr Ser Val Val Lys Gly Ser His Phe Pro
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                            120
Val Gly Val Val Pro Pro Arg Ala Lys Ser Pro Thr Pro Glu Ser Ser
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Thr Ile Ala Ser Tyr Val Thr Leu Arg Lys Thr Lys Lys Met Met Asp
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 Leu Arg Thr Glu Arg Pro Arg Ser Ala Val Glu Gln Leu Cys Leu Ala
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                 165
 Glu Ser Thr Arg Pro Arg Met Thr Val Glu Glu Gln Met Glu Arg Ile
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                                 185
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 Arg Arg Tyr Gln Gln Ala Cys Leu Arg Glu Lys Lys Gly Leu Asn
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 Val Ile Gly Ala Ser Asp Gln Ser Pro Leu Gln Ser Pro
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Cys Glu Gln Asn Leu Leu Ser Arg Pro Asp Gly Ser Ala Ser Phe Leu
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Gln Gly Asp Thr Ser Val Leu Ala Gly Val Tyr Gly Pro Ala Glu Val
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Lys Val Ser Lys Glu Ile Phe Asn Lys Ala Thr Leu Glu Val Ile Leu
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Arg Pro Lys Ile Gly Leu Pro Ala Gly Val Ser Gly Trp Gln Ser Gly
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Glu Lys Ser Arg Glu Arg Leu Ile Arg Asn Thr Cys Glu Ala Val Val
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Val Ser Asp Ala Gly Ser Leu Leu Ala Cys Cys Leu Asn Ala Ala Cys
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Met Ala Leu Val Asp Ala Gly Val Pro Met Arg Ala Leu Phe Cys Gly
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Val Ala Cys Ala Leu Asp Ser Asp Gly Thr Leu Val Leu Asp Pro Thr
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Ser Lys Gln Glu Lys Glu Ala Arg Ala Val Leu Thr Phe Ala Leu Asp
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Ser Val Glu Arg Lys Leu Leu Met Ser Ser Thr Lys Gly Leu Tyr Ser
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Asp Thr Glu Leu Gln Gln Cys Leu Ala Ala Ala Gln Ala Ala Ser Gln
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His Val Phe Arg Phe Tyr Arg Glu Ser Leu Gln Arg Arg Tyr Ser Lys
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Val Leu Ala Val Leu Leu Thr Leu Val Phe Trp Lys Leu Ile Arg Ser
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Arg Arg Ser Ser Gln Arg Ala Val Leu Leu Val Gly Leu Cys Asp Ser
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Gly Lys Thr Leu Leu Phe Val Arg Leu Leu Thr Gly Leu Tyr Arg Asp
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Thr Gln Thr Ser Ile Thr Asp Ser Cys Ala Val Tyr Arg Val Asn Asn
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Asn Arg Gly Asn Ser Leu Thr Leu Ile Asp Leu Pro Gly His Glu Ser
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Val Phe Val Val Asp Ser Ala Ala Phe Gln Arg Glu Val Lys Asp Val
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Ala Glu Phe Leu Tyr Gln Val Leu Ile Asp Ser Met Gly Leu Lys Asn
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Thr Pro Ser Phe Leu Ile Ala Cys Asn Lys Gln Asp Ile Ala Met Ala
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Lys Ser Ala Lys Leu Ile Gln Gln Gln Leu Glu Lys Glu Leu Asn Thr
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Leu Arg Val Thr Arg Ser Ala Ala Pro Ser Thr Leu Asp Ser Ser Ser
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Thr Ala Pro Ala Gln Leu Gly Lys Lys Gly Lys Glu Phe Glu Phe Ser
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Gln Leu Pro Leu Lys Val Glu Phe Leu Glu Cys Ser Ala Lys Gly Gly
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Val Lys Ser His Thr Glu Thr Asp Glu Lys Gln Thr Glu Ser Arg Thr
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Ile Thr Pro Pro Ala Ala Pro Lys Pro Lys Arg Glu Glu Asn Pro Gln
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Glu Tyr Glu Leu Glu Lys Arg Val Glu Arg Leu Glu Leu Phe Pro Val
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Gly Ala Gly Ala Asp Met Gly Leu Glu Lys Leu Gly Ile Phe Val Lys
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Thr Val Thr Glu Gly Gly Ala Ala His Arg Asp Gly Arg Ile Gln Val
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Asn Asp Leu Leu Val Glu Val Asp Gly Thr Ser Leu Val Gly Val Thr
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Gln Ser Phe Ala Ala Ser Val Leu Arg Asn Thr Lys Gly Arg Val Arg
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Phe Met Ile Gly Arg Glu Arg Pro Gly Glu Gln Ser Glu Val Ala Gln
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Leu Ile Gln Gln Thr Leu Glu Gln Glu Arg Trp Gln Arg Glu Met Met
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Arg Lys Leu Gln Ser Leu Glu Gln Glu Lys Gly Arg Trp Arg Val Glu
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Pro Glu Val Gly Asp Leu Leu Arg Asn Lys Leu Val Arg Leu Met Thr
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His Leu Asp Thr Asp Val Lys Arg Val Ala Ala Glu Phe Leu Phe Val
Leu Cys Ser Glu Ser Val Pro Arg Phe Ile Lys Tyr Thr Gly Tyr Gly
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Lys Glu Ala Lys Ala Ser Ile Asn Pro Val Thr Gly Arg Val Glu Glu
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Lys Pro Asp Val Val Gln Asp Lys Glu Thr Glu Arg Asn Leu Gln Arg
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Ile Ala Thr Arg Gly Val Val Gln Leu Phe Asn Ala Val Gln Lys His
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Gln Lys Asn Val Asp Glu Lys Val Lys Glu Ala Gly Ser Ser Met Arg
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Val Leu Arg Gly Met Asp Gly Ser Thr Asn Glu Thr Ala Ser Ser Arg
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 Val Cys Leu His Val Asp Lys Asp Lys Val Ser Val Glu Phe Cys Ser
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705	:				710	)				715	i				Ala 720
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			740	)				745	5				750	,	Phe
		75	5				760	)				765	•		Phe
	770	)				77!	5				780	)			Gly
Pro	o Gly	y Gl	y Ala	a Se			o Arg	g Ala	a Lei	ı Glu	ı Ile	e Asr	Lys	Met	Ile
78!	5			_	790		_ + -··	, n	. <b>-</b> 1.	795		ר (בות	n Met	. Val	800 Val
				80	5				810	)				815	
Tr	p Le	u Pr	o Arg	g Se	r Ala	a Le	u Pro	o Arg	g Alá	a va.	r TT6	AI'S	j n⊥s	3 5 1 (	Asp

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Lys Leu Glu Trp Phe Ser Thr Leu Phe Pro Arg Ile Pro Val Pro Val
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Gln Lys Asn Ile Asp Gln Gln Ile Lys Thr Arg Pro Arg Lys Ile Lys
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Arg Glu Ala Lys Glu Arg Glu Lys Glu Arg Arg Arg Ser Arg Ser Ile
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Asp Arg Gly Leu Glu Arg Arg Arg Ser Arg Ser Arg Glu Arg His Arg
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Ser Arg Ser Arg Ser Arg Asp Arg Lys Gly Asp Arg Asp Arg Asp
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 Tyr Asp Lys Glu Arg Gly Asn Glu Arg Glu Lys Glu Arg Glu Arg Ser
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Ser Leu Ala Tyr Cys Thr Phe Leu Leu Ala Val Gly Leu Ser Arg Ile
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Gln Gln Asp Ala Lys His Ile Leu Glu His Val Phe Phe Gln Val Val
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                          40
                                             45
Gln Glu Ala Gln Ser Leu Gly Gly Gln Cys Val Pro Val Val Cys Asp
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Ser Ser Gln Glu Ser Glu Val Arg Ser Leu Phe Glu Gln Val Asp Arg
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Glu Gln Gln Gly Arg Leu Asp Val Leu Val Asn Asn Ala Tyr Ala Gly
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Val Gln Thr Ile Leu Asn Thr Arg Asn Lys Ala Phe Trp Glu Thr Pro
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          100
Ala Ser Met Trp Asp Asp Ile Asn Asn Val Gly Leu Arg Gly His Tyr
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Phe Cys Ser Val Tyr Gly Ala Arg Leu Met Val Pro Ala Gly Gln Gly
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Leu Ile Val Val Ile Ser Ser Pro Gly Ser Leu Gln Tyr Met Phe Asn
                                     155
                  150
Val Pro Tyr Gly Val Gly Lys Ala Ala Cys Asp Lys Leu Ala Ala Asp
                                  170
               165
Cys Ala His Glu Leu Arg Arg His Gly Val Ser Cys Val Ser Leu Trp
           180 185
Pro Gly Ile Val Gln Thr Glu Leu Leu Lys Glu His Met Ala Lys Glu
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Glu Val Leu Gln Asp Pro Val Leu Lys Gln Phe Lys Ser Ala Phe Ser
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Ser Ala Glu Thr Thr Glu Leu Ser Gly Lys Cys Val Val Ala Leu Ala
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                                      235
Thr Asp Pro Asn Ile Leu Ser Leu Ser Gly Lys Val Leu Pro Ser Cys
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               245
Asp Leu Ala Arg Arg Tyr Gly Leu Arg Asp Val Asp Gly Arg Pro Val
                    265
Gln Asp Tyr Leu Ser Leu Ser Ser Val Leu Ser His Val Ser Gly Leu
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Gly Trp Leu Ala Ser Tyr Leu Pro Ser Phe Leu Arg Val Pro Lys Trp
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                                         300
Ile Ile Ala Leu Tyr Thr Ser Lys Phe
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Val Ser Lys Ser Cys Leu Asp Ser Asp Pro Ala Gly Pro Phe Gln Gly
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Ser Gln Pro Gly Cys His Ser Gly Leu Leu Thr Asn Thr Pro Ala Ala
Leu Val Pro Ala His Ala Arg Gln Arg Ser Gln Pro Ser Leu Leu
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Ser Ser Ser Pro Arg Lys Ser Arg Ser Trp Gln Gly Ser Gly Pro Met
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Trp Pro Gly Pro Gly Tyr Phe Pro Asp Leu Thr Ser Pro Thr Ala Gln
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Pro Leu Gln Leu Leu Gly Ala Leu His Gly Cys Ser Phe Pro Pro
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Leu Pro Ser Gly Gln Pro Cys Pro
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Lys Val Thr Leu Pro Asn Tyr Asp Asn Val Pro Gly Asn Leu Met Leu
                            40
Ser Ala Leu Gly Leu Arg Leu Gly Asp Arg Val Leu Leu Asp Gly Gln
                        55
Lys Thr Gly Thr Leu Arg Phe Cys Gly Thr Thr Glu Phe Ala Ser Gly
                                         75
                    70
Ser Trp Val Gly Val Glu Leu Asp Glu Pro Glu Gly Lys Asn Asp Gly
                                    90
                85
Ser Val Gly Gly Val Arg Tyr Phe Ile Cys Pro Pro Lys Gln Gly Leu
                                105
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Phe Ala Ser Val Ser Lys Ile Ser Lys Ala Val Asp Ala Pro Pro Ser
                                                 125
                            120
 Ser Val Thr Ser Thr Pro Gly Pro Pro Arg Met Asp Phe Ser Arg Val
                                             140
                         135
 Thr Gly Lys Gly Arg Arg Glu His Lys Gly Lys Lys Thr Pro Ser
                                         155
                    150
 Ser Pro Ser Leu Gly Ser Leu Gln Gln Arg Asp Gly Ala Lys Ala Glu
                                     170
                 165
 Val Gly Asp Gln Val Leu Val Ala Gly Gln Lys Gln Gly Ile Val Arg
                                 185
             180
 Phe Tyr Gly Lys Thr Asp Phe Ala Pro Gly Tyr Trp Tyr Gly Ile Glu
                             200
         195
 Leu Asp Gln Pro Thr Gly Lys His Asp Gly Ser Val Phe Gly Val Arg
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                         215
 Tyr Phe Thr Cys Pro Pro Arg His Gly Val Phe Ala Pro Ala Ser Arg
                                         235
                     230
 Ile Gln Arg Ile Gly Gly Ser Thr Asp Ser Pro Gly Asp Ser Val Gly
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250
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Ala Lys Lys Val His Gln Val Thr Met Thr Gln Pro Lys Arg Thr Phe
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Thr Thr Val Arg Thr Pro Lys Asp Ile Ala Ser Glu Asn Ser Ile Ser
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Arg Leu Leu Phe Cys Cys Trp Phe Pro Trp Met Leu Arg Ala Glu Met
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Gln Ser
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120
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Phe Gln Glu Gly Cys Leu Glu Val Gln Trp Gly Gly Arg Gly Phe Gly
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Ser Pro Trp Lys Phe Leu Arg Glu Cys Ser Asn Leu Cys Leu Thr Ile
Met Met Val Val Ser Trp Thr Ala Gly Gly Lys Ala Lys Pro Cys Gly
                        55
Arg Gly Gly Leu Gln Arg Lys Ala Ala Ala Thr Thr Ala Ser Phe
                    70
                                        75
Pro Thr His Ser His Trp Gln Thr Gly Gly Gln Val Gln Ser Pro Lys
                                    90
Glu Thr Ala Ala Cys Ala Gly His Pro Pro Gly Thr Ala Phe Ser Leu
                                105
Ile Leu Pro Val Pro Pro Thr Cys Trp Val Ser Val Ala
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        115
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                                 25
Arg Thr Ala Pro Lys Lys Gln Leu Pro Ser Ile Pro Lys Asn Ala Leu
                                                 45
                             40
Pro Ile Thr Lys Pro Thr Ser Pro Ala Pro Ala Ala Gln Ser Thr Asn
                                             60
Gly Thr His Ala Ser Tyr Gly Pro Phe Tyr Leu Glu Tyr Ser Leu Leu
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Ala Glu Phe Thr Leu Val Val Lys Gln Lys Leu Pro Gly Val Tyr Val
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Gln Pro Ser Tyr Arg Ser Ala Leu Met
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caggccgggc tcaaccaaaa gctgaatttt attgttactg gcttacagga tattgacaag

180

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                                25
Ser Gln Ala Gly Leu Asn Gln Lys Leu Asn Phe Ile Val Thr Gly Leu
                            40
Gln Asp Ile Asp Lys Cys Arg Gln Gln Leu His Asp Ile Thr Val Pro
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                        55
Leu Glu Val Phe Glu Tyr Ile Asp Gln Gly Arg Asn Pro Gln Leu Tyr
Thr Lys Glu Cys Leu Glu Arg Ala Leu Ala Lys Asn Glu Gln Val Lys
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85
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Gly Lys Ile Asp Thr Met Lys Lys Phe Lys Ser Leu Leu Ile Gln Glu
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Leu Ser Lys Val Phe Pro Glu Asp Met Ala Lys Tyr Arg Ser Ile Arg
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Gly Glu Asp His Pro Pro Ser
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Thr Val Glu Gly His Pro Ser Ala Asp Lys Asn Trp Ala Tyr Lys Tyr
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Leu Leu Gly Leu Ile Lys Ser Ser Pro Thr Phe Leu Pro Thr Glu Asp
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Ser Asp Phe Leu Met Ser Leu Asp Leu Glu Ser Arg Asp Gln Ala Trp
                                       315
                   310
Ser Pro Lys Thr Cys Asp Ile Val Ile Glu Gly Ser Gln Ser Pro Thr
                                  330
               325
Ser Pro Ala Ser Ser Ser Pro Lys Pro Gly Ser Lys Ala Gly Ser Gln
                              345
           340
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Trp Gln Glu Val Leu Ser Asp Ser Gln Arg Glu His Leu Gln Gln Phe
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Leu	Pro	Gln	Phe		Glu	Asp	Ser	Ala 105	Glu	Gln	Gln	Asn	Glu 110	Leu	Ile
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		195					200			Thr		205			
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_				245					250	Ser				255	
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				325					330	Lys				335	
	_		340					345		Thr			350		
		355					360			Pro		365			
	370					375					380				Ser
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				405					410					415	Pro
			420					425					430		Glu
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465					470					475					Gln 480
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			500					505					510		Gln
Glu	Asn	Glu	Asp	ser	ser	Asp	АТа	ınr	ınr	Pro	val	Pro	arg	٧d⊥	Arg

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_		595			Ala		600					605			
	610				Pro	615					620				
Glu	Leu	Leu	Lys	Asp	Ser	Gln	Phe	Leu	Ala	Pro	Asp	Val	Thr	Ser	
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				645	Val				650					655	
-	_		660		Lys			665					670		
		675			Ser		680					685			
	690				Lys	695					700				
	Pro	Ser	Lys	Val	Lys	Ser	Ser	Ser	Lys	Glu 715	Ser	Ser	Ile	Lys	Val 720
705	C	C ~ ~	<i>α</i> 1	Dwa	710 Ser	Glu	Gln	Sar	Gln		Ser	Len	Ser	Asp	
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		755			Ile		760					765			
	770				Val	775					780				
Leu 785	Leu	Val	Ser	Ser	Pro 790		Met	Pro	His	Leu 795	Gly	Thr	Met	Leu	ser 800
Pro	Ala	Ser	Ser	Gln	Thr		Pro	ser	Ser		Ala	Ala	Ala	Arg	Val
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			820					825					830		Ala
		835			Ala		840					845			
Thr	Leu 850		Gln	Met	Pro	Ala 855	Gly	Pro	Gln	Ile	Arg 860		Pro	Ala	Thr
Ala	Thr	Gln	Thr	Lys	Val	Val	Pro	Gln	Thr	Val	Met	Ala	Thr	Val	Pro
865					870					875					880
				885					890					895	Gly
			900					905					910		Val
Ser	Lys	Pro 915		Thr	Ser	Ser	Pro 920		Thr	Ser	Ala	Pro 925		Ala	Ser
Thr	Ala 930	Ala		Ile	Gln	Asn 935		Thr	Gly	Gln	Asn 940		Ile	Lys	Gln
Val			Thr	Gly	Gln			Val	Lys	Pro			Gly	Asn	Ser

945		950		955	960			
Ile Pro Leu	Thr Ala '	Thr Asn Phe	Arg Ile 970	Gln Gly Lys	Asp Val Leu 975			
Arg Leu Pro	Pro Ser :	Ser Ile Thr	Thr Asp .	Ala Lys Gly	Gln Thr Val 990			
Leu Arg Ile 995	Thr Pro	Asp Met Met 1000		Leu Ala Lys 1009				
Thr Thr Val	Lys Leu '	Thr Gln Asp 1015	Leu Phe	Gly Thr Gly 1020	Gly Asn Thr			
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His Ala Ala	Asp Ser 1045	Pro Ala Lys	Ala Ser 1050		Ala Pro Ser 1055			
Ser Thr Pro	Thr Gly '	Thr Thr Val	Val Lys 1065	Val Thr Pro	Asp Leu Lys 1070			
Pro Thr Glu 107		Ser Ser Ala 1080		Leu Met Pro 1089				
Val Ser Val 1090	Ala Asp	Gln Lys Gly 1095	Lys Ser	Thr Val Ala 110 <b>0</b>	Ser Ser Glu			
Ala Lys Pro 1105		Thr Ile Arg 1110		Gln Gly Leu 1115	Gly Val Met 1120			
Pro Pro Lys	Ala Gly 0	Gln Thr Ile	Thr Val		Ala Lys Gln 1135			
Gly Ala Ser	Val Ala : 1140	Ser Gly Ser	Gly Thr	Val His Thr	Ser Ala Val 1150			
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Gly Lys Leu	Pro Thr 1	Arg Ile Thr	Val Pro 1210		Ile Ser Gln 1215			
Pro Met Lys	Gly Lys : 1220	Ser Val Val	Thr Ala 1225	Pro Ile Ile	Lys Gly Asn 1230			
Leu Gly Ala 123		Ser Gly Leu 1240		Asn Ile Ile 1249				
Met Pro Ala 1250	Gly Thr	Lys Leu Ile 1255	Ala Gly	Asn Lys Pro 1260	Val Ser Phe			
Leu Thr Ala 1265		Leu Gln Gln 1270		Gln Gln Gly 1275	Gln Ala Thr 1280			
Gln Val Arg	Ile Gln 1285	Thr Val Pro	Ala Ser 1290		Gln Gly Thr 1295			
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ggggatatgg 180	ttcttttaga	acctctcaat	gaggagacct	tcatcaacaa	cctcaagaag
cgctttgacc 240	acagtgaaat	atacacttac	attggaagtg	tggttatatc	tgttaaccca
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tatgaactga 360	gccctcacat	ctttgccctt	tcggatgaag	catacagatc	cctacgagat
caagataagg 420	accaatgtat	tctcattact	ggggaaagtg	gagcaggaaa	aacagaggcc
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gactacttca 1440	ataatgctat	catttgtgac	ctaatagaaa	ataacacaaa	tggaatcctg
gccatgttgg 1500	atgaagagtg	cctcagacct	ggcacagtca	ctgatgagac	cttcttagaa
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Asn Asn Leu Lys Lys Arg Phe Asp His Ser Glu Ile Tyr Thr Tyr Ile
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Gly Ser Val Val Ile Ser Val Asn Pro Tyr Arg Ser Leu Pro Ile Tyr
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                                         60
Ser Pro Glu Lys Val Glu Glu Tyr Arg Asn Arg Asn Phe Tyr Glu Leu
                                      75
                  70
Ser Pro His Ile Phe Ala Leu Ser Asp Glu Ala Tyr Arg Ser Leu Arg
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Asp Gln Asp Lys Asp Gln Cys Ile Leu Ile Thr Gly Glu Ser Gly Ala
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Gly Lys Thr Glu Ala Ser Lys Leu Val Met Ser Tyr Val Ala Ala Val
                         120
Cys Gly Lys Gly Ala Glu Val Asn Gln Val Lys Glu Gln Leu Leu Gln
                                         140
                      135
Ser Asn Pro Val Leu Glu Ala Phe Gly Asn Ala Lys Thr Val Arg Asn
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                                     155
Asp Asn Ser Ser Arg Phe Gly Lys Tyr Met Asp Ile Glu Phe Asp Phe
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                                  170
Lys Gly Asp Pro Leu Gly Gly Val Ile Ser Asn Tyr Leu Leu Glu Lys
          180 185
Ser Arg Val Val Lys Gln Pro Arg Gly Glu Arg Asn Phe His Val Phe
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                         200
Tyr Gln Leu Leu Ser Gly Ala Ser Glu Glu Leu Leu Asn Lys Leu Lys
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Leu Glu Arg Asp Phe Ser Arg Tyr Asn Tyr Leu Ser Leu Asp Ser Ala
                                      235
Lys Val Asn Gly Val Asp Asp Ala Ala Asn Phe Arg Thr Val Arg Asn
Ala Met Gln Ile Val Gly Phe Met Asp His Glu Ala Glu Ser Val Leu
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Ala Val Val Ala Ala Val Leu Lys Leu Gly Asn Ile Glu Phe Lys Pro
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Glu Leu Lys Glu Ile Cys Glu Leu Thr Gly Ile Asp Gln Ser Val Leu
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                                      315
Glu Arg Ala Phe Ser Phe Arg Thr Val Glu Ala Lys Gln Glu Lys Val
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Ser Thr Thr Leu Asn Val Ala Gln Ala Tyr Tyr Ala Arg Asp Ala Leu
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Ala Lys Asn Leu Tyr Ser Arg Leu Phe Ser Trp Leu Val Asn Arg Ile
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Asn Glu Ser Ile Lys Ala Gln Thr Lys Val Arg Lys Lys Val Met Gly
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Val Leu Asp Ile Tyr Gly Phe Glu Ile Phe Glu Asp Asn Ser Phe Glu
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Gln Phe Ile Ile Asn Tyr Cys Asn Glu Lys Leu Gln Gln Ile Phe Ile
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                                    410
Glu Leu Thr Leu Lys Glu Glu Glu Glu Tyr Ile Arg Glu Asp Ile
                                                    430
                                425
            420
Glu Trp Thr His Ile Asp Tyr Phe Asn Asn Ala Ile Ile Cys Asp Leu
                            440
Ile Glu Asn Asn Thr Asn Gly Ile Leu Ala Met Leu Asp Glu Glu Cys
                        455
Leu Arg Pro Gly Thr Val Thr Asp Glu Thr Phe Leu Glu Lys Leu Asn
                                        475
                    470
Gln Val Cys Ala Thr His Gln His Phe Glu Ser Arg Met Ser Lys Cys
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Pro His Phe Asn Lys His Leu Leu Gly Ala Glu His Gly Asp Glu Pro
Arg His Gly Gly Leu Thr Leu Arg Leu Gly Leu His Gln Gln Ser Val
Leu Gly Gly Gln Asp Gln Leu Arg Val Arg Val Thr Glu Leu Glu Asp
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His Leu Ala Val Asp Gly Asp Arg Ala Ala Trp Pro Val Gly Ile
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                                          60
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Pro Ala Pro Ser Arg Pro Ala Ser Arg Phe Glu Val Leu Arg Trp Asp
                                       75
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Tyr Phe Thr Glu Gln His Ala Phe Ser Cys Ala Asp Gly Ser Pro Arg
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Thr Ala Leu Glu Glu Leu Asn Arg Arg Tyr His Pro Ala Leu Arg Leu
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Gln Lys Gln Gln Leu Val Asn Gly Tyr Arg Arg Phe Asp Pro Ala Arg
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Gly Met Glu Tyr Thr Leu Asp Leu Gln Leu Glu Ala Leu Thr Pro Gln
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                                       155
Gly Gly Arg Arg Pro Leu Thr Arg Arg Val Gln Leu Leu Arg Pro Leu
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                                   170
Ser Arg Val Glu Ile Leu Pro Val Pro Tyr Val Thr Glu Ala Ser Arg
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           180
Leu Thr Val Leu Leu Pro Leu Ala Ala Glu Arg Asp Leu Ala Pro
                           200
Gly Phe Leu Glu Ala Phe Ala Thr Ala Ala Leu Glu Pro Gly Asp Ala
                       215
                                           220
Ala Ala Ala Leu Thr Leu Leu Leu Tyr Glu Pro Arg Gln Ala Gln
                                       235
Arg Val Ala His Ala Asp Val Phe Ala Pro Val Lys Ala His Val Ala
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                                   250
Glu Leu Glu Arg Arg Phe Pro Gly Ala Arg Val Pro Trp Leu Ser Val
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Gln Thr Ala Ala Pro Ser Pro Leu Arg Leu Met Asp Leu Leu Ser Lys
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Lys His Pro Leu Asp Thr Leu Phe Leu Leu Ala Gly Pro Asp Thr Val
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                                           300
Leu Thr Pro Asp Phe Leu Asn Arg Cys Arg Met His Ala Ile Ser Gly
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Trp Gln Ala Phe Phe Pro Met His Phe Gln Ala Phe His Pro Ala Val
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qaccetqaca ceacaqaagt caatttgaac aacattqaqa acatcacaac acagaccett

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                            40
Ser Asp Asn Ser Lys Pro Lys Ile Phe Lys Ser Gln Ile Glu Asn Ile
                        55
                                            60
Asn Leu Thr Asn Gly Ser Asn Gly Arg Asn Thr Glu Ser Pro Ala Ala
                                        75
Ile His Pro Cys Gly Asn Pro Thr Val Ile Glu Asp Ala Leu Asp Lys
                                    90
Ile Lys Ser Asn Asp Pro Asp Thr Thr Glu Val Asn Leu Asn Asn Ile
                                105
Glu Asn Ile Thr Thr Gln Thr Leu Thr Arg Phe Ala Glu Ala Leu Lys
                            120
Asp Asn Thr Val Val Lys Thr Phe Ser Leu Ala Asn Thr His Ala Asp
                        135
                                            140
Asp Ser Ala Ala Met Ala Ile Ala Glu Met Leu Lys Val Asn Glu His
                    150
                                        155
Ile Thr Asn Val Asn Val Glu Ser Asn Phe Ile Thr Gly Lys Gly Ile
                                    170
                                                        175
Leu Ala Ile Met Arg Ala Leu Gln His Asn Thr Val Leu Thr Glu Leu
                                                    190
            180
                                185
Arg Phe His Asn Gln Arg His Ile Met Gly Ser Gln Val Glu Met Glu
                            200
                                                205
Ile Val Lys Leu Leu Lys Glu Asn Thr Thr Leu Leu Arg Leu Gly Tyr
                        215
His Phe Glu Leu Pro Gly Pro Arg Met Ser Met Thr Ser Ile Leu Thr
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235
225
                    230
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7	т1 о	Mot	uio			Glu	Gl n	Hic			Wa l	Acn	Pro		Tyr
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			T		mh.	T10			ת דת	Thr	Gl n		Ser	Met	Ara	
Met		Gln	ьeu	ser	ınr			val	M±d	TITE	940	TEU	561		3	
_	930		-	D1-	7	935		C1	D~~	ጥኤ⊶		T1 ***	Tle	Δen	Asn	
_		Asp	Leu	Phe			тте	GIU	Pro			TAL	TIG	vah	Asp 960	
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Glu	Asp	Ile	Val	Asn	Gln	Glu	Thr			Val	Ala	Ser			Leu	
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Gln Asp Ile				car	Ara				Lvs	Ͳν·	Ara	Asn	Ile
GIN Asp IIe			PIO	DCI		1065		7124	_,_	-1-	1070		
	1060	_		<b>~</b> 1				T10	Dvo	Lou			Val
Leu Ser Ser	Gln	Ser .	Met				TTE	TIE	PIO			PIO	Val
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1105			1110		-			1115					1120
Arg Gln Val	นาไ				Ser	Δla	Asn	Met	Asp	Pro	Ala	Met	Met
Arg Gin vai				1111	001		1130		<u>-</u> -			1135	;
		1125			a1				7 an	C . ~	7 cn		
Phe Arg Gln			ьeu	Ser	GIH			TILL	ASII	Ser			шси
	1140					1145		_		_	1150		_
Asp Val Glr	Gly	Gly	Ala	His	Lys	Lys	Arg	Ala	Arg	Arg	ser	ser	Leu
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	шец		1190			. •		1195	;			•	1200
1185 Gln Met Met	0				G1	D×a				Thr	Leu	Thr	
GIn Met Met	ser			пр	GIU	PIO			СТУ	ILL	пси	1219	= 5,5
		1205				_	1210		_	~7			
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	1220					1225					1230		_
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1250				1255					1260				
Pro Ile Arg	Tire	T	<b>a</b> 1			T	7 ~~	Desa	717	Lan	λen	Thr	Ser
				(4)7			ASD	Pro	Ala				
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Lys Tyr Glu Asp Val Glu His Lys Val Thr Thr Val Phe Gly Gln Pro
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ggggtagaag 2520	tctaacccct	ccaccccctc	tcctccccag	cagtcccacg	cgggtatggg
agagaatgaa 2580	gttctttgtc	tctaagggat	tcaaaccaga	aacggaggga	cctctggttc
ccagagggag 2640	gaaaatccat	gatgtctgct	gcccagggag	ctattgccac	cgcctccttg
ggatgaagta 2700	ttgccagcta	ccaacagttc	cttcccaacg	gccatcttcc	agccttctta
aacgactcct 2760	agcatcttcg	ggaggctcct	gaaggactga	agcaaaggaa	atctctgaag
ggatttagtc 2820	cttgaaaggg	agtagggata	cttagggtgt	tctgtgttga	gcgcttcttc
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<210> 4858

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<211> 269
<212> PRT
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Gly Trp Trp Arg Leu Gly Ser Ser Ser Gln Ala Ala Cys Leu Lys Gln
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Ile Leu Leu Gln Leu Asp Leu Ile Glu Gln Gln Gln Gln Leu
                           40
Gln Ala Lys Glu Lys Glu Ile Glu Glu Leu Lys Ser Glu Arg Asp Thr
                       5.5
Leu Leu Ala Arg Ile Glu Arg Met Glu Arg Met Gln Leu Val Lys
                   70
                                       75
Lys Asp Asn Glu Lys Glu Arg His Lys Leu Phe Gln Gly Tyr Glu Thr
               85
                                   90
Glu Glu Arg Glu Glu Thr Glu Leu Ser Glu Lys Ile Lys Leu Glu Cys
           100
                              105
Gln Pro Glu Leu Ser Glu Thr Ser Gln Thr Leu Pro Pro Lys Pro Phe
                                              125
                           120
Ser Cys Gly Arg Ser Gly Lys Gly His Lys Arg Lys Ser Pro Phe Gly
                                          140
                       135
Ser Thr Glu Arg Lys Thr Pro Val Lys Lys Leu Ala Pro Glu Phe Ser
                   150
                                       155
Lys Val Lys Thr Lys Thr Pro Lys His Ser Pro Ile Lys Glu Glu Pro
                                   170
               165
Cys Gly Ser Leu Ser Glu Thr Val Cys Lys Arg Glu Leu Arg Ser Gln
                                                  190
                               185
Glu Thr Pro Glu Lys Pro Arg Ser Ser Val Asp Thr Pro Pro Arg Leu
                          200
Ser Thr Pro Gln Lys Gly Pro Ser Thr His Pro Lys Glu Lys Ala Phe
                                          220
                       215
Ser Ser Glu Ile Glu Asp Leu Pro Tyr Leu Ser Thr Thr Glu Met Tyr
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Leu Cys Arg Trp His Gln Pro Pro Pro Ser Pro Leu Pro Leu Arg Glu
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Ser Ser Pro Lys Lys Glu Glu Thr Val Ala Ser Lys Ala
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<211> 689
<212> DNA
<213> Homo sapiens
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ggcctcccac ggtgcctgtg ctgggtggcg gtggtggtgc caagaggaat ggaatgtcct
gggctccttc aggagctctc tacccagggg caaggagagc ccagagagaa gcgccctggt
240
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ctettgaget teetgatetg etectgteec eegeteteet eeacteeett geettteeet
aggttqtccc ctccctgggc ttttgtgtgt tttgggagat gtcacctaac caggacattg
360
atattcaatc ccatccccct tcctcccacc ctgccccact ttgatttaat cctttggctg
tgggctgagg cctcccaggg aagttgggtg gggtgggtgt tgagaccccc tcagaccagc
acagagacct gtccttgtgc agtctgcacc ctgcactccc tcccttgcct gtagatgttc
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689
<210> 4860
<211> 173
<212> PRT
<213> Homo sapiens
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Met Arg Thr Arg Leu Phe Ala Val Pro Gly Arg Val Ala Lys Glu Asp
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Trp Thr Leu Asp Leu Glu Pro Arg Gly Pro Val His Ile His Pro Thr
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            20
Arg Val Ser Gly Gly Leu Pro Arg Cys Leu Cys Trp Val Ala Val Val
                                                 45
                            40
Val Pro Arg Gly Met Glu Cys Pro Gly Leu Leu Gln Glu Leu Ser Thr
                        55
Gln Gly Gln Gly Glu Pro Arg Glu Lys Arg Pro Gly Leu Leu Ser Phe
                                        75
                    70
Leu Ile Cys Ser Cys Pro Pro Leu Ser Ser Thr Pro Leu Pro Phe Pro
                                    90
                85
Arg Leu Ser Pro Pro Trp Ala Phe Val Cys Phe Gly Arg Cys His Leu
                                105
            100
Thr Arq Thr Leu Ile Phe Asn Pro Ile Pro Leu Pro Pro Thr Leu Pro
                            120
His Phe Asp Leu Ile Leu Trp Leu Trp Ala Glu Ala Ser Gln Gly Ser
                        135
                                            140
Trp Val Gly Trp Val Leu Arg Pro Pro Gln Thr Ser Thr Glu Thr Cys
                                                             160
                    150
                                        155
Pro Cys Ala Val Cys Thr Leu His Ser Leu Pro Cys Leu
                165
<210> 4861
<211> 1622
<212> DNA
<213> Homo sapiens
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60
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	tgcggcccgg	ccttcgggtg	ttagactgtg	gggcagctcc	tggggcctgg
	cggtgcagaa	ggtcaacgcc	gcaggcacag	atcccagctc	teetgttgge
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	tgattctgag	cgacatggcg	cccaatgcca	cagggttccg	ggacctcgat
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	cattcctttg	taaaacctgg	gctggaagtc	aaageegteg	gttacagagg
	aggaattcca	gaatgtaagg	atcatcaaac	ctgaagccag	caggaaagag
	tgtacttctt	ggccacacag	taccacggaa	ggaagggcac	tgtgaagcag
	ttgtgccatt	ttcataatgg	tcattagctc	cttttaagct	agaaacgtag
cctgagetee 900	tgaagagttc	ctgggagatt	tgagctgatt	ttggagatgg	agcaggacaa
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ctaagttcag 1020	gggccatgga	aaatgaaaaa	gtccgctata	ttgtgatttg	ggaagagaaa
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	ttaaactgtc	acacacaaac	aggctttcca	cccctgctct	gagagcacca
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tttctggtga 1500	acaatatagc	aattcacgca	ttcttcaagc	agcaaaagtt	cccggaacaa
	cgtatggtct	gaatttatcc	aggcagtggg	tetgetttgg	tttttgctgg
	cagtgtctgg	gctcccaaga	acataaatgt	aattgccaaa	gcaaaaaaaa

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<210> 4862
<211> 260
<212> PRT
<213> Homo sapiens
<400> 4862
Leu Gln Thr Ser Gly Gly Ala Leu Gln Ala Arg Gly Thr Pro Met Ala
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Gly Tyr Leu Lys Leu Val Cys Val Ser Phe Gln Arg Gln Gly Phe His
                              25
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Thr Val Gly Ser Arg Cys Lys Asn Arg Thr Gly Ala Glu His Leu Trp
                          40
Leu Thr Arg His Leu Arg Asp Pro Phe Val Lys Ala Ala Lys Val Glu
                      55
Ser Tyr Arg Cys Arg Ser Ala Phe Lys Leu Leu Glu Val Asn Glu Arg
                  70
                                     75
His Gln Ile Leu Arg Pro Gly Leu Arg Val Leu Asp Cys Gly Ala Ala
              85
                                 90
Pro Gly Ala Trp Ser Gln Val Ala Val Gln Lys Val Asn Ala Ala Gly
          100
                             105
Thr Asp Pro Ser Ser Pro Val Gly Phe Val Leu Gly Val Asp Leu Leu
                         120
                                             125
His Ile Phe Pro Leu Glu Gly Ala Thr Phe Leu Cys Pro Ala Asp Val
                      135
                                         140
Thr Asp Pro Arg Thr Ser Gln Arg Ile Leu Glu Val Leu Pro Gly Arg
                  150
                                     155
Arg Ala Asp Val Ile Leu Ser Asp Met Ala Pro Asn Ala Thr Gly Phe
                                 170 175
              165
Arg Asp Leu Asp His Asp Arg Leu Ile Ser Leu Cys Leu Thr Leu Leu
                             185
Ser Val Thr Pro Asp Ile Leu Gln Pro Gly Gly Thr Phe Leu Cys Lys
                         200
Thr Trp Ala Gly Ser Gln Ser Arg Arg Leu Gln Arg Arg Leu Thr Glu
                     215
                                         220
Glu Phe Gln Asn Val Arg Ile Ile Lys Pro Glu Ala Ser Arg Lys Glu
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                                    235
Ser Ser Glu Val Tyr Phe Leu Ala Thr.Gln Tyr His Gly Arg Lys Gly
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Thr Val Lys Gln
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<210> 4863
<211> 355
<212> DNA
<213> Homo sapiens
<400> 4863
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accatcaacc ctgaggacga cacggatcct ggccatgctg acctggtcct ctatatcact
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aggtttgacc tggagttgcc tgatggtaac neggcagtgc ggggegtcac ccagetgggc
ggggcctgct ccccaacctg gagctgcctc attaccgagg acactggctt cgacctggga
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355
<210> 4864
<211> 118
<212> PRT
<213> Homo sapiens
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Leu Gly Ala His Phe Arg Val His Leu Val Lys Met Val Ile Leu Thr
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Glu Pro Glu Gly Ala Pro Asn Ile Thr Ala Asn Leu Thr Ser Ser Leu
                                25
Leu Ser Val Cys Gly Trp Ser Gln Thr Ile Asn Pro Glu Asp Asp Thr
                            40
Asp Pro Gly His Ala Asp Leu Val Leu Tyr Ile Thr Arg Phe Asp Leu
                        55
                                            60
Glu Leu Pro Asp Gly Asn Xaa Ala Val Arg Gly Val Thr Gln Leu Gly
                    70
                                        75
Gly Ala Cys Ser Pro Thr Trp Ser Cys Leu Ile Thr Glu Asp Thr Gly
                                    90
                85
Phe Asp Leu Gly Val Thr Ile Ala His Glu Ile Gly His Ser Phe Gly
                                105
                                                     110
Leu Glu His Asp Gly Ala
        115
<210> 4865
<211> 444
<212> DNA
<213> Homo sapiens
<400> 4865
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aaggoottog oogacagoto ttacotgott ogocaccago goactcacto tggccagaag
ccctacaagt gcccacattg tggcaaggcc ttcggcgaca gctcctacct cctgcgacac
caqcqcaccc acaqccacga gcggccctac agctgcaccg agtgcggcaa gtgctatagc
300
cagaactcgt ccctgcgcag ccatcagagg gtgcacaccg gtcagaggcc cttcagctgt
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420
gecegggaga agecetteae gegt
444
<210> 4866
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4048

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<211> 148
<212> PRT
<213> Homo sapiens
<400> 4866
Thr Gly Glu Lys Pro Tyr Lys Cys Glu Val Cys Ser Lys Ala Phe Ser
Gln Ser Ser Asp Leu Ile Lys His Gln Arg Thr His Thr Gly Glu Arg
                                25
            20
Pro Tyr Lys Cys Pro Arg Cys Gly Lys Ala Phe Ala Asp Ser Ser Tyr
                            40
Leu Leu Arg His Gln Arg Thr His Ser Gly Gln Lys Pro Tyr Lys Cys
                        55
Pro His Cys Gly Lys Ala Phe Gly Asp Ser Ser Tyr Leu Leu Arg His
                                        75
Gln Arg Thr His Ser His Glu Arg Pro Tyr Ser Cys Thr Glu Cys Gly
                                    90
Lys Cys Tyr Ser Gln Asn Ser Ser Leu Arg Ser His Gln Arg Val His
            100
                                105
Thr Gly Gln Arg Pro Phe Ser Cys Gly Ile Cys Gly Lys Ser Phe Ser
                            120
Gln Arg Ser Ala Leu Ile Pro His Ala Arg Ser His Ala Arg Glu Lys
   130
                       135
                                            140
Pro Phe Thr Arg
145
<210> 4867
<211> 391
<212> DNA
<213> Homo sapiens
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cettetecae atececatte tggtaggaaa agteacecat gecaggatat eeccageeca
gagacagece cagggggtge tgeetggaga cageegggat agetteagte teetgaceet
gacacgggct gcaccaccag acaatgggca ttttcaggcc agactctggc acaaagagaa
ggggcagggc caaggctatg gcccacaagc tcctcagcag ctgagatggg tgcaggaggt
agegetetae teccataget ecceaetgta t
391
<210> 4868
<211> 125
<212> PRT
<213> Homo sapiens
<400> 4868
Met Gly Val Glu Arg Tyr Leu Leu His Pro Ser Gln Leu Leu Arg Ser
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10
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Leu Trp Ala Ile Ala Leu Ala Leu Pro Leu Leu Phe Val Pro Glu Ser
            20
                                25
Gly Leu Lys Met Pro Ile Val Trp Trp Cys Ser Pro Cys Gln Gly Gln
                            40
Glu Thr Glu Ala Ile Pro Ala Val Ser Arg Gln His Pro Leu Gly Leu
                                            60
                        55
Ser Leu Gly Trp Gly Tyr Pro Gly Met Gly Asp Phe Ser Tyr Gln Asn
                                        75
                    70
Gly Asp Val Glu Lys Glu Ala Asp Val Pro Arg Leu Val Ala Ser Phe
                                    90
Cys Pro Ser His Pro Pro Thr Lys Asp Met Arg Leu Leu Pro Ser Asn
                                105
Leu Leu Gly Ala Ser Pro Asp Arg Thr Pro Ser Gly Ile
                            120
<210> 4869
<211> 418
<212> DNA
<213> Homo sapiens
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caqqactqca cggactgcct ggggaggggt ctttggcccc ccggttcctg caggggggct
cggggaggcc ctgtgagcag ttggtcacag gtgggtccca ttcgatgcga tcctgttcct
ccccaacaqc cctggagaag ggggacgttg cctgctgtgg ctgcggctgt tttcctggcc
tgtgagaggc ggggccagag tggccgttgg gaatctgggt gttgcaaggt gaccacaaac
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<210> 4870
<211> 125
<212> PRT
<213> Homo sapiens
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Met Ala Met Gly Ile Gly Trp Glu Leu Asn Gly Val Ala Thr Phe Gly
Trp Thr Arg Arg Gln Pro Ser Phe Leu Gly Gln Asp Cys Thr Asp Cys
Leu Gly Arg Gly Leu Trp Pro Pro Gly Ser Cys Arg Gly Ala Arg Gly
Gly Pro Val Ser Ser Trp Ser Gln Val Gly Pro Ile Arg Cys Asp Pro
                        55
Val Pro Pro Gln Gln Pro Trp Arg Arg Gly Thr Leu Pro Ala Val Ala
                    70
                                        75
Ala Ala Val Phe Leu Ala Cys Glu Arg Arg Gly Gln Ser Gly Arg Trp
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95
                                    90
                85
Glu Ser Gly Cys Cys Lys Val Thr Thr Asn Ser Ser Leu Gly Glu Glu
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Glu Glu Asn Ala Ile Asp Phe Gln Glu Pro Ser Glu Val
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                                                125
        115
<210> 4871
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<212> DNA
<213> Homo sapiens
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caqcccctca qqccatqctg ctgctcagct gcatggcaaa gtcctgcaca tgctccttca
180
gagtetggeg ggeatetgee tgtgeeeget tetecegtge eegeteetge tgeagettgg
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540
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caageegget agaetteeeg teeteeeett eeegaetgea tteagteeeg eegggaeegt
1260
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tecqetteac eteccaceca caggiteaag cetecteagt atetgagaaa ggegegaage
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<210> 4872
<211> 90
<212> PRT
<213> Homo sapiens
<400> 4872
Gly Arg Lys Arg Leu Gln Ser Cys Trp Ala Ala Pro Arg Ser Val Gln
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Gln Pro Leu Arg Pro Cys Cys Cys Ser Ala Ala Trp Gln Ser Pro Ala
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His Ala Pro Ser Glu Ser Gly Gly His Leu Pro Val Pro Ala Ser Pro
                            40
                                                45
Val Pro Ala Pro Ala Ala Ala Trp Ser Val Ser Thr Ala Ala Ala Ala
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Pro Ala Ala Cys Arg Pro Ala Ala Gly Ala Gly Pro Cys Gln Gly His
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                                        75
Gln Gly Leu Pro Gly Ser Pro Leu Pro Glu
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<210> 4873
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<212> DNA
<213> Homo sapiens
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cacagococa tocaggitos catggiggos ggotococto toaggacaas coagaigtgo
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600
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720
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cgaacacatg gcatcctgcc aggatgacct gaagtcatcc tcacctttcc tttccacata
aagceggeee atacacettt tetttggaae taacecacea gatettagaa gatgtacaeg
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<211> 128
<212> PRT
<213> Homo sapiens
<400> 4874
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Asp Leu Ser Pro Asp His Pro Gly Thr Glu Leu Trp Asp Ser Val Val
                                                    30
Leu Glu Asn His Val Val Thr Asp Glu Asp Glu Pro Ala Leu Lys Arg
                            40
Gln Arg Leu Glu Ile Asn Cys Gln Asp Pro Ser Ile Lys Ser Phe Leu
                                            60
Tyr Ser Ile Asn Gln Thr Ile Cys Leu Arg Leu Asp Ser Ile Glu Ala
                    70
Lys Leu Gln Ala Leu Glu Ala Thr Cys Lys Ser Leu Glu Glu Lys Leu
                85
                                    90
Asp Leu Val Thr Asn Lys Gln His Ser Pro Ile Gln Val Pro Met Val
                                105
Ala Gly Ser Pro Leu Arg Thr Thr Gln Met Cys Asn Lys Val Arg Trp
                                                 125
                            120
        115
<210> 4875
<211> 1255
<212> DNA
<213> Homo sapiens
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480
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Phe Met Asn Ile Arg Leu Ala Lys Val Thr Tyr Thr Asp Arg Trp Gly
His Gln Val Lys Leu Asp Asp Leu Phe Val Thr Gly Arg Asn Val Arg
Tyr Val His Ile Pro Asp Asp Val Asn Ile Thr Ser Thr Ile Glu Gln
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Thr Thr Val Leu Phe Trp Gly Phe Ser Lys Ala Ser Pro Val Val Leu
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Arg Gly His Ser Glu Gln Ala Asn Thr Ala Arg Val Thr His Tyr Thr
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Gln Arg Lys Asp Asn Glu Gln Met Ala Ile Val Glu Asn Ser Val Val
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Cys Phe Ser Asn Ala Thr Tyr Phe Ser Arg Gln Val Ile Leu Pro Met
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Cys His Arg Pro Glu His Arg Thr Val Ile Met Gln Arg Ala Val Thr
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Lys Ser Leu Asp Val Leu Asp Leu His Gly Asn Gln Ile Thr Lys Ile
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Asn Phe Leu Ser His Val Asp Asn Leu Asn Gly Leu Asp Ser Leu Thr
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Glu Leu Asn Leu Arg His Asn Gln Ile Thr Phe Val Arg Asp Val Asp
             150 155
Asn Leu Pro Cys Leu Gln His Leu Phe Leu Ser Phe Asn Asn Ile Ser
    165 170
                              175
Ser Phe Asp Ser Val Ser Cys Leu Ala Asp Ser Ser Ser Leu Ser Asp
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Ile Thr Phe Asp Gly Asn Pro Ile Ala Gln Glu Ser Trp Tyr Lys His
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Thr Val Leu Gln Asn Met Met Gln Leu Arg Gln Leu Asp Met Lys Arg
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Glu Glu Lys Lys Arg Glu Ser His Lys Gln Ser Leu Leu Lys Glu Lys
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Val Asn Phe Thr Arg Xaa Glu Trp Arg Glu Leu Asp Leu Ala Gln Arg
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Ala Lys Ala Ala Ile Val Cys Tyr Asp Leu Thr Asp Ser Ser Ser Phe
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Glu Arg Ala Lys Phe Trp Val Lys Glu Leu Arg Ser Leu Glu Glu Gly
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Cys Gln Ile Tyr Leu Cys Gly Thr Lys Ser Asp Leu Leu Glu Glu Asp
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Arg Arg Arg Arg Val Asp Phe His Asp Val Gln Asp Tyr Ala Asp
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4077

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Val Asp Glu Cys Ser Leu Ala Glu Lys Thr Cys Val Arg Lys Asn Glu
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Glu Gln Lys Gln Gln Pro Pro Asn Ser Phe Ser Gln Gln His Ser Glu
Thr Gln Gly Ala Glu Lys Pro Asp Pro Glu Ser Ser His Ser Pro Pro
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Arg Tyr Thr Asp Gln Gly Gly Glu Glu Glu Asp Tyr Glu Ser Glu
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                                    90
Glu Gln Leu Gln His Arg Ile Leu Thr Ala Ala Leu Glu Phe Val Pro
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Ala His Gly Trp Thr Ala Glu Ala Ile Ala Glu Gly Ala Gln Ser Leu
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Gly Leu Ser Ser Ala Ala Ala Ser Met Phe Gly Arg Met Gly Ser Glu
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Leu Ile Leu His Phe Val Thr Gln Cys Asn Thr Arg Leu Thr Arg Val
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Leu Ile Pro Tyr Ile Glu His Trp Pro Arg Ala Leu Ser Ile Leu Met
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Leu Pro His Asn Ile Pro Ser Ser Leu Ser Leu Leu Thr Ser Met Val
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Asp Asp Met Trp His Tyr Ala Gly Asp Gln Ser Thr Asp Phe Asn Trp
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Tyr Thr Arg Arg Ala Met Leu Ala Ala Ile Tyr Asn Thr Thr Glu Leu
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Val Met Met Gln Asp Ser Ser Pro Asp Phe Glu Asp Thr Trp Arg Phe
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Val Gly Arg Ser Gln Arg Gly Pro Thr Pro Gln Asn Ala His Lys Ser
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Trp Asn Gln Leu Val Thr Ala Ala Gly Pro Ser Arg Pro Ile Trp Ile
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Lys Ser Ser Xaa Gly Gly Thr His Gly Ile Leu Gly Gly His Leu Arg
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Ala Pro Pro Pro Thr Ile Pro Pro Ser Lys Val Ala Ser Glu Cys Glu
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Gly Arg Gly Lys Gln Thr Pro Ala Pro His Ser Pro Ser Leu Pro His
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Ala Ser Ile Leu Arg Trp Pro Glu Ala Leu Pro Leu Arg Gln Ile Met
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Thr Pro Asp Ala Ser Ser Pro Leu Tyr Pro Cys His Met Glu Gly Pro
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Lys His Leu Ala Leu Asn Cys Lys Trp Lys Pro Pro Gln Pro Leu His
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			820					825		Leu		_	830	_	_
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Glu Gly Ala Arg Pro Gly Gly Ile Ile His Val Tyr Gly Asp Asp Ser
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Asn Ser Arq Ala Leu Gly Val Met Asp Lys Ser Thr Ala Ile Pro Lys
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Ala Ser Ser Ser Glu Ser Leu Ser Ala Lys Thr Cys Ser Leu Phe Leu
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Pro Asn Tyr Val Gln Asp Lys Tyr Leu Leu Gln Leu Leu Arg Asn Ala
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Asp Asp Val Ser Thr Trp Val Ala Ala Glu Ile Val Thr Ser His Thr
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Ser Lys Leu Gln Val Asn Leu Leu Ser Lys Phe Xaa Leu Ile Ala Lys
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Phe Leu Lys Ser Asp Ser Leu Cys Leu Met Glu Gly Arg Arg Phe Arg
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His	Leu		Tyr	Glu	Leu	Gln		Ala	ьуѕ	мет	ьeu	Thr 205	Cys	TIE	ASII
_	_	195	5		m	C 0 20	200	Cln.	Glu	Dro	Tle	Cys	Ser	Ala	Pro
Ala		Lys	Pro	HIS	пр	215	Ser	GIII	Giu	110	220	0,0			
<b>G</b>	210	01	ת 1 ת	37 n T	uic		Δla	Thr	Tle	Glv		Val	Leu	Ser	Pro
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Gly	Leu	Leu	Ser	Glu		Asn	Thr	Ile	Arg	lle	GIU	Pne	THE	Ser	Asp 320
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Gln	Ala	Arg	Ala		Ser	Thr	Pne	ASI	330		PHE	Giu	AIG	335	Glu
_			<b>~</b>	325	~1	Dree	П	т1д			Glv	Asn	Phe		Thr
Lys	GIY	HIS	340		GIU	PLO	IYI	345		ADII	<b>U</b>		350		
Cor	7 cm	Dro	ምክ _ዮ	ጥህዮ	Asn	Tle	Glv			Val	Glu	Phe	Thr	Cys	Asp
Ser	Asp	355		1 <b>y</b> 1			360					365			
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Trp	Pro	Glu			Val	Glu	. Gly			Cys	Ile	Trp	vsv гъх	тте	His
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Val	Gly			Lys	Arg	Ile			Asp	, тте	: GII	445	; 115u	. ADI	Leu
_		435	) . 7	. 71~		ጥሎ~	440 - Tle		- Agr	Glu	, Agr			Met	Pro
Ser			Asp	, тте	ьeu	455		. <u> </u>	. Ast	. Gry	460	)			
u: -	450 - ⊤1	T.A	י נים	r (dlm	. ጥህዮ			Ast	Ser	Glv			Lys	Leu	Tyr
#15		. הפו	. Giy	<b>U1</b> 1	470		. <b></b> 1			475			-		480
Ser	, - Ser	Th:	Pro	Ast			: Ile	Glr	n Phe			Asp	Pro	Ala	Gly
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570

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565

545

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Lys Ile Phe Leu Pro Lys Lys Leu Leu Glu Cys Leu Pro Arg Cys Pro
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Leu Leu Pro Pro Glu Arg Leu Arg Trp Asn Thr Asn Glu Glu Ile Ala
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_			180					Asn 185					190		
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Lys Tyr Lys Gln Leu Thr Trp Ile Ala Leu Lys Phe Ala Leu Tyr Lys
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Pro Leu Glu Ala Lys Gly Leu Ala Thr Gln Gly Ala Ser Leu Pro Leu
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Leu Pro Thr Val Thr Cys Val Ser Ile Lys Ser Trp Lys Met Glu Cys
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Thr Asn Cys Pro Pro Lys Glu Gln Pro Gly Asp Leu Phe Asn Glu Asp
Trp Asp Ser Glu Leu Lys Ala Asp Gln Gly Asn Pro Tyr Asp Ala Asp
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Asp Ile Gln Glu Ser Ile Ser Gln Glu Leu Lys Pro Trp Val Cys
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Ala Pro Gln Gly Asp Met Ile Tyr Asp Pro Ser Trp His His Pro Pro
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Val Met Asp Gly Val Ile Ser Asp His Glu Cys Gln Glu Leu Gln Arg
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Leu Thr Asn Val Ala Ala Thr Ser Gly Asp Gly Tyr Arg Gly Gln Thr
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Ser Pro His Thr Pro Asn Glu Lys Phe Tyr Gly Val Thr Val Phe Lys
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Ala Leu Lys Leu Gly Gln Glu Gly Lys Val Pro Leu Gln Ser Ala His
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Leu Tyr Tyr Asn Val Thr Glu Lys Val Arg Arg Ile Met Glu Ser Tyr
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Phe Arg Leu Asp Thr Pro Leu Tyr Phe Ser Tyr Ser His Leu Val Cys
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395

390

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170

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120

115

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Pro Ser Gly Lys Ser Val Gly Glu Ala His Ser Val Ser Pro Pro Pro
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His Val Asp Cys Ala Arg Ala Arg Pro Thr Gly Ser Cys Thr Pro Glu
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Gln Gln Gly Ile Leu Glu Lys Glu Leu Leu Val Arg Tyr Leu Glu Gln
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Cys Pro Thr Thr Ser Gly Thr Asp Phe Pro Ser Leu Gln Ser Lys Ala
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Gly Leu Ile Ser Val Asn Ser Gly Ala Pro Ala Ser His Glu Cys Ala
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gatgcattta 780	tgaaagcaaa	tectggetae	aaatggtgtc	ctaccacaaa	caagcctgtg
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Val Leu Asp Pro Lys Glu Lys Gln Lys Tyr Thr Asp Met Ala Lys Glu
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Tyr Lys Asp Ala Phe Met Lys Ala Asn Pro Gly Tyr Lys Trp Cys Pro
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Thr Thr Asn Lys Pro Val Lys Ser Pro His Pro Leu Ser Ile His Glu
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Arg Asn Phe Gly Pro Ser His Leu Thr Leu Gln Glu Thr Cys Gln Ala
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Val Ile Leu Ile Phe Cys Leu Met Thr Leu Ile Gly Asn Leu Phe Ile
Ile Ile Leu Thr Tyr Leu Asp Ser His Leu His Thr Pro Leu Tyr Phe
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Phe Leu Ser Asn Leu Ser Phe Leu Asp Leu Cys Tyr Thr Thr Ser Ser
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Ile Pro Gln Leu Leu Val Ser Leu Trp Gly Val Glu Lys Thr Ile Ser
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Tyr Ala Gly Cys Met Val Gln Leu Tyr Phe Phe Leu Thr Leu Gly Thr
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Thr Glu Cys Val Leu Leu Val Val Met Ser Tyr Asp Arg Tyr Ala Ala
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Val Cys Arg Pro Leu His Tyr Thr Val Leu Met His Ser Arg Phe Cys
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His Leu Leu Ala Val Ala Ser Trp Val Ser Gly Phe Thr Asn Pro Ala
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Ala Gly Pro Gln Arg Val Leu Pro Gly Glu Arg Glu Glu Arg Pro Pro
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Ala Thr His Val Tyr Arg Tyr His Arg Gly Glu Ser Lys Leu His Met
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4209

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1500

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Gln Ala Ser Ser Asn Ala Pro Gly Ala Pro Ala Gln Gln Trp Leu Thr
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Gln Val Thr Cys Thr Pro Gly Pro Ala Leu Pro Ala Arg His Ser Pro
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Pro Ser Ala Leu Ser Gly Gly Gln Pro Ala Asp Thr Gln Thr Arg Ala
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Thr Ser Lys Ser Leu Leu Pro Val Arg Ser Lys Glu Val Asp Val Ser
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Arg Arg Asn Val Arg Lys Gly Tyr Lys Pro Leu Ser Lys Gln Lys Ser
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Thr Gln Lys Val Glu Leu Leu Glu Lys Phe Arg Asp Asn Cys Leu Ala
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Asp Met Val Ala Cys Cys Leu Phe Ser Cys Ser Ser Lys His Tyr Pro
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Glu Tyr Cys Phe Thr Arg Lys Glu Gly Leu Ser Lys Cys Gly Arg Cys
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Met His Lys Leu Glu Cys Ser Pro Met Val Val Phe Gly Glu Asn Trp
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Cys Asn Gly Phe Thr Ile Glu Asp Glu Glu Leu Ser His Leu Gly Ser
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Ala Ile Phe Pro Asp Val Ala Leu Met Asn His Ser Cys Cys Pro Asn
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Val Ile Val Thr Tyr Lys Gly Thr Leu Ala Glu Val Arg Ala Val Gln
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Glu Ile Lys Pro Gly Glu Glu Val Phe Thr Ser Tyr Ile Asp Leu Leu
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Tyr Pro Thr Glu Asp Arg Asn Asp Arg Leu Arg Asp Ser Tyr Phe Phe
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Thr Cys Glu Cys Gln Glu Cys Thr Thr Lys Asp Lys Asp Lys Ala Lys
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Val Glu Ile Arg Lys Leu Ser Asp Pro Pro Lys Ala Glu Ala Ile Arg
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Asp Met Val Arg Tyr Ala Arg Asn Val Ile Glu Glu Phe Arg Arg Ala
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Gln Glu Lys Met Ser Ser Val Phe Glu Asp Ser Asn Val Tyr Met Leu
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Val Gln Leu Cys Thr Pro Leu Leu Pro Arg Asn Arg Gln Ile Tyr
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Pro Arg Lys Leu Asp Arg Phe His Pro Lys Glu Leu Leu Glu Cys Ala
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465	_	<b>.</b>	<b>01.</b>		470	Sor	. Acn	Δla	Ser	475 Glv		Phe	Glu	Ile	His
Asn	Leu	ггуя	GT.	485		Jer	LDII	1114	490	)				495	
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		515	;				520					525	•		Leu
Glr	Thr		Lev	ı Ala	a Asr	Leu 535		Gln	Lys	Ser	Ser 540	: Glu	ı Ile	Glu	Ile
Leu	ı Ala	, a Val	L Ası	. Lei	ı Pro			Thr	: Ile	e Leu	Gln	Phe	e Lev	Ser	Leu
545	5				550	)				555	5				560
Glu	ı Tr	) Ası	) Ala			ı Glr	n Ala	Phe	Asr 570	ı Thr	Thr	· Val	Lys	Glr 579	Leu
T 01	1 601	- Dro	ים, די	569 1 Pro		s Glr	ı Arc	TVI			Leu	ı Val	L Cys		Glu
			58	0				585	5				590	)	
Ile	е Ту		ı Ile	e Lys	s Val	i Glu	1 Lys		s Val	l Ser	(Va)	L Let 60!	ı Phe	e Lei	ı Tyr
0	л Пээ <b>э</b>	59! 220		ם אפו	יט <b>י"ו</b> ים	r Tvi			e Lei	ı Phe	<b>=</b>	J J .	-		
sei	т т.Х.	ר ייידי	ים ה	ادت م	1 -	1.		,							

615

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465	;				470					475	5				His 480
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